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OM protein - protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Scoring table:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Title:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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Pred. No. 1s the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  A_Geneseq_101002:*
1: /SIDS1/gcgdata/ge
2: /SIDS1/gcgdata/ge
3: /SIDS1/gcgdata/ge
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      March 24, 2003, 17:16:57; Search time 25.6875 Seconds (without alignments) 46.686 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    BLOSUM62
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                                                                                                                                       | SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1980.DAT:*
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1981.DAT:*
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1982.DAT:*
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1983.DAT:*
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1983.DAT:*
| SIDS1/gcgdata/geneseq/geneseqp-emb1/AA1984.DAT:*
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SUMMARIES

1 2 3 3 4 4 6 6 7 7 10	Result
41 41 41 41 41 41	Score
993333322	Query Match I
21 25 25 31 31 33 33 33 142 208	Query Match Length DB
14 AAR38001 14 AAR38005 14 AAR38005 14 AAR36797 14 AAR36799 11 AAR35079 21 AAB15054 21 AAB15058 21 AAB15058 21 AAB17686	ID
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ALIGNMENTS

RESULT 1 AAR38001

AAR38001 standard; protein; 21

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PD XX PR XX PD XX PR XX PR XX PR XX PR XX XX PR XX XX XX XX
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                                                                                                                                                                                                                                                                                                                                              AAR38001;
                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                             Prion protein region B #5.
                                                                                                                                                                                                                                                                                                                      14-OCT-1993 (first entry)
                                                                                                                                                                                                                       resistance.
                                                                                    03-DEC-1991;
10-JUL-1992;
                                                                                                                                                                        WO9311155-A.
          WPI; 1993-196994/24.
                                                         (PROT-) PROTEUS MOLECULAR DESIGN LTD.
                                                                                                                       03-DEC-1992;
                                                                                                                                                10-JUN-1993.
                                  Fishleigh RV, Mee RP,
                                                                                    91GB-0025747.
92GB-0014663.
                                                                                                                          92WO-GB02246.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
W09311155-A
                                         Misc-difference
                                                                       Misc-difference 24
                                                                                                     Misc-difference
                                                                                                                                  Misc-difference
                                                                                                                                                                 Misc-difference
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                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                       Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                            Antigen; prion; protein; region; frame shift; repeat; mutation; prpc; FSa; FSb; subfragment; antibody; treatment; spongiform encephalopathy; human; sheep; cattle; cellular binding; aggregation; mammal; scraple; immune system; prpsc; ratio-inverso peptide; enzymatic degradation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prion protein region B subfragment #1.
                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                   Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 14-OCT-1993 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAR38003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR38003 standard; protein; 25 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 10; Page 65; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          e.g. Creutzfeld-Jacob disease and kuru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New polypeptide(s) contg. antigenic site of prion protein - useful for treatment and diagnosis of mammalian encephalopathies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       11 GSDYEDRYY 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
les 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            21 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                     /note= "May be absent"
23
                /note= "One or more residue or may be absent"
                                                                                                                                                                                                                                                             /note= "May be absent"
                                                                                /note= "May be absent"
                                                                                                                                                                                                                                                                                                                                  Location/Qualifiers
                                                     'note- "May be absent"
                                                                                                                                            note= "May be absent"
                                                                                                                                                                           'note= "May be absent"
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                                                                                                                                                                                                                                                                                            /note= "One or more residues or may be absent"
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77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
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Pred. No. 0.22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 14; Length 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         2; Indels
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RESULT 3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ΡŢ
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                                   Misc-difference
                                                                                   Misc-difference
                                                                                                                                   Misc-difference
                                                                                                                                                                                                             Synthetic
                                                                                                                                                                                                                                                             Antigen; prion; protein; region; frame shift; repeat; mutation; prpc; FSa; FSb; subfragment; antibody; treatment; spongiform encephalopathy; human; sheep; cattle; cellular binding; aggregation; mammal; scrapie; immune system; prpsc; ratio-inverso peptide; enzymatic degradation;
                                                                                                                                                                                                                                                          resistance.
                                                                                                                                                                                                                                                                                                                                                                                            Prion protein region B subfragment #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          AAR38005 standard; protein; 25
                                                                                                                                                                                                                                                                                                                                                                                                                                                      14-0CT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAR38005;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           e.g. Creutzfeld-Jacob disease and kuru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New polypeptide(s) contg. antigenic site of prion protein - useful for treatment and diagnosis of mammalian encephalopathies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 1993-196994/24.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PROT-) PROTEUS MOLECULAR DESIGN LTD.
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10-JUL-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                              (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Mee RP,
       /note= "May be absent"
                                                                                                                                                     Location/Qualifiers
                                                       /note= "May be absent"
                                                                                           /note- "One or more residues or may be absent"
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92GB-0014663.
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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Misc-difference

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AAR36797
ID AARX
XX AAR
AC AAR
XX 14-0
DT 14-0
XX Pr1-
XX Ant
                                                                                                                                                                                                                                                                                                                                              The sequences given in AAR38002-05 represent polypeptide subfragments CC derived from an antigenic site, region B, of a prion protein. Prion CC proteins comprise six regions of interest (A-F), and two related CC proteins having a nucleic acid coding sequence frame shift mutation in CC region E having a nucleic acid coding sequence frame shift mutation CC (FSa) or 1 (FSb). These subfragments or the full length peptide CC (see AAR36796-99 and AAR38000-01), and antibodies raised against these, CC (see AAR36796-99 and AAR38000-01), and antibodies raised against these, CC (see AAR36796-99 and protein specific mercephalopathy in humans, CC may be used to treat or prevent spongiform encephalopathy in humans, CC sheep or cattle. They can be used to block cellular binding and CC system. These peptides may be used to distinguish between the normal CC system. These peptides may be used to distinguish between the normal CC ratio-inverso peptide modification to improve resistance to enzymatic CC decrafation protein (PrPC) and the scrapic associated form (PrPSc).
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                                                                                                                                               RESULT 4
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Misc-difference 24
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10-JUL-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     New polypeptide(s) contg. antigenic site of prion protein useful for treatment and diagnosis of mammalian encephalopathies
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fishleigh RV,
                                                                                                                                                                                                                                                                                                             Sequence
                                                                                       AAR36797;
                                                            14-OCT-1993
                                                                                                                   AAR36797 standard; protein; 31 AA
Antigen; prion; protein; region; frame shift; repeat; mutation; PrPc;
                             Prion protein region B #1.
                                                                                                                                                                                            12 GSDYEDRYY 20
                                                                                                                                                                                                                                                    Local Similarity
les 7; Conserv
                                                                                                                                                                                                                     1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                             25 AA;
                                                                                                                                                                                                                                                    Conservative
                                                            (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           91GB-0025747.
92GB-0014663.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "May be absent" 22
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "May be absent"
5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "One or more residue or may be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "May be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note= "May be absent"
                                                                                                                                                                                                                                                                   93.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Robson B;
                                                                                                                                                                                                                                                       0
                                                                                                                                                                                                                                                                   Score 41; L
Pred. No. 0.
                                                                                                                                                                                                                                                        Mismatches
                                                                                                                                                                                                                                                                                   DB 14; Length 25;
                                                                                                                                                                                                                                                        2
                                                                                                                                                                                                                                                          0,
                                                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                             0
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Matches Query Match

Local Similarity 77.1 pes 7; Conservative

93.28;

Score 41; DB 1 Pred. No. 0.33; Mismatches

DB 14; Length 31;

Indels

0

Gaps

0

0

1 GXDXEDRYY 9 _ _ _ _ _

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FSa; FSb; subfragment; antibody; treatment; spongiform encephalopathy; human; sheep; cattle; cellular binding; aggregation; mammal; scraple; immune system; PrPsc; ratio-inverso peptide; enzymatic degradation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "May be absent" Misc-difference 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "May be absent" Misc-difference 27
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                resistance.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference 31
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 /note= "May be absent" Misc-difference 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
                          polypeptides derived from an antigenic site, region B, of a prion protein. Prion proteins comprise six regions of interest (A-F), and two related frame shift peptides sequences caused by a repeating section in region B having a nucleic acid coding sequence frame shift mutation of +1 (FSa) or -1 (FSb). These peptides and subfragments of these (see AAR38002-05), and antibodies raised against these, may be used to treat or prevent spongiform encephalopathy in humans, sheep or cattle. They can be used to block cellular binding and aggregation of prion proteins and to stimulate the mammalian immune system. These peptides may be used to distinguish between the normal form of prion protein (PrPc) and the scrapic-associated form (PrPsc). These peptides may include rare or synthetic amino acids or a ratio-inverso peptide modification to improve resistance to enzymatic degradation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        03-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                      03-DEC-1991;
10-JUL-1992;
                                                                                                                                                                                                                                The sequences given in AAR36797-99 and AAR38000-01 represent
                                                                                                                                                                                                                                                                   Claim 9; Page 64; 82pp; English.
                                                                                                                                                                                                                                                                                                 e.g. Creutzfeld-Jacob disease and kuru
                                                                                                                                                                                                                                                                                                              New polypeptide(s) conty. antigenic site of prion protein useful for treatment and diagnosis of mammalian encephalopathies
                                                                                                                                                                                                                                                                                                                                                              WPI; 1993-196994/24.
                                                                                                                                                                                                                                                                                                                                                                                           Fishleigh RV,
                                                                                                                                                                                                                                                                                                                                                                                                                         (PROT-) PROTEUS MOLECULAR DESIGN LTD
Sequence
 31 AA;
                                                                                                                                                                                                                                                                                                                                                                                              Mee RP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                          91GB-0025747.
92GB-0014663.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        92WO-GB02246.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   note= "One or more residues or may be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note- "One or more residue or may be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "May be absent"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              /note= "May be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             'note- "May be absent"
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GSDYEDRYY 20

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The sequences given in AAR36797-99 and AAR36000-01 represent polypeptides derived from an antigenic site, region B, of a prion protein. Prion proteins comprise six regions of interest (A-F), and tweeleast frame shift peptides sequences caused by a repeating section in region E having a nucleic acid coding sequence frame shift mutation of +1 (FSa) or -1 (FSb). These peptides and subfragments of these (see AAR38002-05), and antibodies raised against these, may be used to treat or prevent spongiform encephalopathy in humans, sheep or cattle. They can be used to block cellular binding and aggregation of prion
                                                                                                                                            Claim 9; Page 65; 82pp; English.
                                                                                                                                                                     e.g. Creutzfeld-Jacob disease and kuru
                                                                                                                                                                               New polypeptide(s) contg. antigenic site of prion protein useful for treatment and diagnosis of mammalian encephalopa
                                                                                                                                                                                                                                                        Fishleigh RV, Mee RP,
                                                                                                                                                                                                                                                                                                                                                   03-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                (PROT-) PROTEUS MOLECULAR DESIGN LTD
                                                                                                                                                                                                                                                                                                            03-DEC-1991;
10-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                             10-JUN-1993.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "May be absent"
Misc-difference 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Antigen; prion; protein; region; frame shift; repeat; mutation; PrPc; FSa; FSb; subfragment; antibody; treatment; spongiform encephalopathy; human; sheep; cattle; cellular binding; aggregation; mammal; scrapie; immune system; PrPsc; ratio-inverso peptide; enzymatic degradation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      resistance
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Prion protein region B #3.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         14-OCT-1993 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAR36799 standard; protein; 31 AA.
                                                                                                                                                                                                                                                                                                           91GB-0025747.
92GB-0014663.
                                                                                                                                                                                                                                                                                                                                                 92WO-GB02246
                                                                                                                                                                                                                                                                                                                                                                                                                                            /note= "May be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          /note= "May be absent" 28
                                                                                                                                                                                                                                                                                                                                                                                                                                 /note-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  'note- "May be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note- "May be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "May be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "One or more residues or may be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            note- "May be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         /note= "May be absent"
                                                                                                                                                                                                                                                                                                                                                                                                                       "One or more residue or may be absent"
                                                                                                                                                                                                                                                     Robson B;
                                                                                                                                                                                 encephalopathies
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through a linker to a solid phase, forming a complex witch its coupled where (A) comprises 10-50 amino acids capable of forming a secondary comprises the (A)-solid phase. The presents a secondary comprises an immune mediator coupled on (A). The present sequence compresents a peptide used in an example from the production of commination derivatives, characterised by covalently attaching molecules at the incorporation into an immunestimulating complex (Iscom) resulting an examplex complex can be used as a scaffold for the production of the incorporation into an immunostimulating complex (Iscom) resulting an example from the present sequence complex which is used for the chemical coupling of antigenic cone or more peptides having fibronectin. (A) derivatised with cone or more peptides having fibronectin. laminin- or vitronectin. like binding activities can be used for the promotion of cell-attachment to plastic surfaces, in particular to inhibit tumour growth and metastasis, complex is such diagnostic (A) molecules could be used to detect molecules could for or indicative of pregnancy or as a diagnostic derived from or indicative of pregnancy or of a disease, such as an example could disease, such as an
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        밁
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មិន្ត្រីស្ត្រីនេះ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AAW35545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
infectious, autoimmune or cancerous disease
                                                                                                                                                                                                                                                                                                                                                                      A non-dendritic peptide carrier (A) has been developed which is coupled through a linker to a solid phase, forming a complex of (A) solid phase.
                                                                                                                                                                                                                                                                                                                                                                                                                    Example 31; Page 156; 262pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         diagnostic agent and as a scaffold for production of chemical
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Non-dendritic peptide carrier linked to a solid phase - useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Heegaard PMH,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             03-APR-1996;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     scaffold; inhibition; metastasis; wound healing; solid phase
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (PEPR-) PEPRESEARCH AS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 03-APR-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      16-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W09738011-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Synthetic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     T-cell stimulatory peptide; immunogen; non-dendritic; carrier; tumour;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Immunization DNDPC SEQ ID NO:89 from WO9738011.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                22-APR-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW35545 standard; peptide; 33 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                proteins and to stimulate the mammalian immune system. These peptidemay be used to distinguish between the normal form of prion protein (PrPc) and the scrapic-associated form (PrPsc). These peptides may include rare or synthetic amino acids or a ratio-inverso peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      AAW35545;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                modification to improve resistance to enzymatic degradation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    12 GSDYEDRYY 20
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
les 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       31 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Jakobsen PH;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       96DK-0000398
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             97WO-DE00146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 41; |
Pred. No. 0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 14; Length 31;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       These peptides
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0,
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RRESULT 7
AAB15054
ID AAB15
XX AAB1
AC AAB1
XX AAB1
XX AB1
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Best Local Similarity
Thes 7; Conserv
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                                                                                                                                                       Query Match
Best Local
                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                        The present invention relates to a method for reducing the risk of scoring a false positive test result in testing a sample for aberrant prion protein. The method involves the use of guandine thiocyanate (gdnSCN) or its functional equivalent. This test is highly useful for testing for transmissible spongiform encephalopathies (TSEs) such as BSE (bovine spongiform encephalopathy). The method allows a faster, simpler and more reliable method for monitoring cattle and sheep for the presence of aberrant prion protein before it reaches the human and animal food chain. In the invention antipeptide antibodies were raised against sheep prion protein peptides. The present sequence is the human prion protein sequence homologous to the sheep peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Use of guanidine thiocyanate for reducing risk of false-positive results in testing mammalian sample for aberrant prion protein, useful for detection of transmissible spongiform encephalopathies -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Prion; PrP; guanidine thiocyanate; gdnSCN; TSE; BSE;
transmissible spongiform encephalopathy; antibody;
bovine spongiform encephalopathy; sheep; cattle; human; mouse;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-DEC-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB15054 standard; Peptide;
                                                                                                                                                                                                                                              Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 2; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2000-506099/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Schreuder BEC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-FEB-2000; 2000WO-NL00079.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             hamster; rabbit.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Human prion protein peptide homologous to ovine sequence 145-177.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (DIEN-) STICHTING DIENST LANDBOUWKUNDIG ONDERZOE.
                                                                                                                    Local Similarity
bes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 GXDXEDRYY 9
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                                                          1 GXDXEDRYY 9
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                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Jacobs JG,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      $
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bossers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93.2%;
                                                                                                                                                 93.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Langeveld JPM,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ω
ω
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Pred. No. 0.
                                                                                                                                                 Score 41; DB 2
Pred. No. 0.36;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 18;
                                                                                                                                                                                 DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        .36;
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                                                                                                                       ?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 33;
                                                                                                                                                                                 Length 33;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Van Keulen LJM;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0
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RESULT 8
AAB15058
ID AAB1
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AAW17686
ID AAW1
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XXEXEXXXXXX
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                                                                                                                                                                                                                                                                                                                                                Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                     scoring a false positive test result in testing a sample for aberrant prion protein. The method involves the use of guanddine thiocyanate (gdnSCN) or its functional equivalent. This test is highly useful for testing for transmissible spongiform encephalopathles (TSEs) such as BSE (bovine spongiform encephalopathy). The method allows a faster, simpler and more reliable method for monitoring cattle and sheep for the presence of aberrant prion protein before it reaches the human and animal food chain. In the invention antipeptide antibodies were raised against sheep prion protein peptides. The present sequence is the cattle prion protein sequence homologous to the sheep peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB15058 standard; Peptide; 33 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Use of guanidine thiocyanate for reducing risk of false-positive results in testing mammalian sample for aberrant prion protein, useful for detection of transmissible spongiform encephalopathies -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Prion; PrP; guanidine thiocyanate; gdnSCN; TSE; BSE; transmissible spongiform encephalopathy; antibody; bovine spongiform encephalopathy; sheep; cattle; hum
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              18-DEC-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cattle prion protein peptide homologous to ovine sequence 145-177.
                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               The present invention relates to a method for reducing the risk of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Disclosure; Fig 2; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Schreuder BEC,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Garssen GJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-FEB-2000; 2000WO-NL00079
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200048003-A1
                                Prion protein peptide Hu 90-231.
                                                                                                                                             AAW17686 standard;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (DIEN-) STICHTING DIENST LANDBOUWKUNDIG ONDERZOE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         17-AUG-2000
                                                                       14-JAN-1998
                                                                                                                                                                                                                                                                                                               Local Similarity
nes 7; Conserv
                                                                                                                                                                                                                                        1 GSDYEDRYY 9
                                                                                                                                                                                                                                                                          1 GXDXEDRYY 9
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                                                                                                                                                                                                                                                                                                             Conservative
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                                                                     (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bossers
                                                                                                                                             peptide; 142 AA.
                                                                                                                                                                                                                                                                                                                               93.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Langeveld JPM,
                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                               Score 41; DB 2
Pred. No. 0.36;
                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                    DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Smits MA,
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                                                                                                                                                                                                                                                                                                                                                  Length 33;
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PrP; alpha helical domain; screening; inhibition;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                            WO200029850-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         The present sequence represents a prion protein (PrP) peptide. PrP has an ability to induce a conformational change in cellular prion protein (PrP-c). Methods, for screening compounds which inhibit the binding of PrP-c to a PrP peptide, are used for screening of drugs that may be useful in the treatment prion-related disease e.g. scrapie, BSE (bovine spongiform encephalopathy), CJD (Creutzfeldt-Jakob disease), kuru, GSS (Gerstmann-Straussler-Scheinker disease) and FFI (fatal familial insomnia).
                                                                                                                                          Modified-site
                                                                                                                                                                  Disulfide-bond
                                                                                                                                                                                                                                                   Region
                                                                                                                                                                                                                                                                                                                                 Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human prion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAB07318
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-OCT-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAB07318 standard; protein; 208 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 11; Page 7-38; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Prion proteins (PrPs) having at least one alpha-helical domain - used in assays for screening compounds able to inhibit or decreate the binding of PrP peptide(s) to cellular prion proteins or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        binding; scrapie; bovine spongiform encephalopathy; BSE; CJD; Creutzfeldt-Jakob disease; kuru; GSS; FFT; fatal familial insomnia; Gerstmann-Straussler-Scheinker disease; hamster; human.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WPI; 1997-272248/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cohen FE,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      02-NOV-1995;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  peptide(s)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (REGC ) UNIV CALIFORNIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28-OCT-1996;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                53 GSDYEDRYY 61
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                               prion protein; transmissible spongiform encephalopathy;
spongiform encephalopathy; BSE diagnosis; TSE; prp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    142 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Kaneko K,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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                                                                                                                                                      /note= "Repeat region consisting of tandem repeats
of repeat unit: PHGGGWGQ (AAB07319)"
157...192
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                                                                     /note= "C-terminal phospho-inositol glycolipid
membrane anchor (-GPI)"
                                                                                                                                          208
                                                                                                                                                                                                                                                Location/Qualifiers 29..69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93.28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 41; |
Pred. No. 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 18;
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AAB07329
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       The present sequence is the human prion protein (PrP) sequence. Conversion of the normal cellular form of PrP into an aggregated, insoluble isoform is implicated in the pathogenesis of Transmissible Spongiform Encephalopathies (TSEs). Examples of TSEs include Bovine Spongiform Encephalopathy (BSE), scraple, Creutzfeldt-Jakob disease (CJD) and Gerstmann-Straussler-Sheinker syndrome (GSS). The concentration of this protein in body fillid or tissue samples may be measured by an assay of the present invention, in which a PrP ppitope is captured by an antibody, which is then detected. The presence of PrP indicates TSE. PrP epitopes (AAB07320-B07326) are derived from the protease resistant core of PrP that is occluded when the PrP is in an aggregated state.
                                     27-OCT-1999;
                                                                 25-MAY-2000.
                                                                                        WO200029849-A1
                                                                                                                                           Modified-site
                                                                                                                                                         Disulfide-bond
                                                                                                                                                                                                                                     Homo sapiens.
                                                                                                                                                                                                                                                                            Human;
                                                                                                                                                                                                                                                                                                   Human prion protein sequence.
                                                                                                                                                                                                                                                                                                                                                                               AAB07329 standard; protein; 208 AA
                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                             Human; prion protein; transmissible spongiform encephalopathy; bovine spongiform encephalopathy; TSE diagnosis; PrP.
                                                                                                                                                                                                                                                                                                                               17-OCT-2000
                                                                                                                                                                                                                                                                                                                                                           AAB07329;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Disclosure; Page 43-44; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Novel immunoassay for prion protein, used for the determination of transmissible spongliform encephalopathies in bovines {\tt -}
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hope J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                  120 GSDYEDRYY 128
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(BBSR-) BBSRC C
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                27-OCT-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     1 Similarity
7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Barnard GJR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          208 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                              (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99WO-FI00897
                                                                                                                                              208
                                                                                                                                                                /note= "Repeat region consisting of tandem repeats
of repeat unit: PHGGGWGQ (AAB07319)"
                                                                                                                                                                                                Location/Qualifiers 29..69
                                                                                                                 /note= "C-terminal phospho-inositol glycolipid
membrane anchor (-GPI)"
                                                                                                                                                        157..192
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   93.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Birkett CR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Score 41;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 21; Length 208; 2.5;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Indels
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17-NOV-1998;

98FI-0002480 99WO-FI00896.

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RESULT 12
AAB07317
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The present sequence is the human prion protein (PrP) sequence. Conversion of the normal cellular form of PrP into an aggregated, insoluble isoform is implicated in the pathogenesis of Transmissible Spongiform Encephalopathies (TSES). Examples of TSES include Bovine Spongiform Encephalopathy (BSE), scrapie, Creutzfeldt-Jakob disease (CJD) and Gerstmann-Straussler-Sheinker syndrome (GSS). The concentration of this protein in body fluid or tissue samples may be measured by an assay of the present invention, in which a PrP pilope is captured by an antibody, which is then detected. The presence of PrP indicates TSE PrP epitopes (AAB07320-B07326) are derived from the protease resistant core of PrP that is occluded when the PrP is in an aggregated state.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Disclosure; Page 43-44; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New immunoassay for prion protein, used for determination of transmissible spongiform encephalopathies in mammals, comprises
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hope J, Barnard GJR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 specific capture antibody
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                                                                                                                                                                                                                                                                                                                                                                                                                      Cattle prion protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAB07317;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB07317 standard; protein; 217 AA.
                                                                                                                                                                                                                                                                                                                                                                       Cattle; prion protein; transmissible spongiform encephalopathy; bovine spongiform encephalopathy; BSE diagnosis; TSE; PrP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                    17-OCT-2000 (first entry)
                                                                                                                                                                               WO200029850-A1
                                                                                                                                                                                                                                       Disulfide-bond Modified-site
                                                                                                                                                                                                                                                                                                Region
                                                                                                                                                                                                                                                                                                                                              Bos bovis.
              Hope J,
                                                                                        17-NOV-1998;
                                                                                                                    27-OCT-1999;
                                                                                                                                                 25-MAY-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       120 GSDYEDRYY 128
                                          (WALL-) WALLAC OY.
(BBSR-) BBSRC OFFICE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local Similarity
nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GXDXEDRYY 9
              Barnard GJR,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            208 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                      98FI-0002481
                                                                                                                    99WO-FI00897.
                                                                                                                                                                                                                                                                    /note= "Repeat region consisting of tandem repeats
of repeat unit: PHGGGWGQ (AAB07319)"
                                                                                                                                                                                                                                                                                                  Location/Qualifiers 37..79
                                                                                                                                                                                                         /note= "C-terminal phospho-inositol glycolipid
membrane anchor (-GPI)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 93.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Birkett CR;
               Birkett CR:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 41; DB 21; Length 208; Pred. No. 2.5;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     2:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is the cattle prion protein (PrP) sequence. Conversion of the normal cellular form of PrP into an aggregated, insoluble isoform is implicated in the pathogenesis of Transmissible Spongiform Encephalopathies (TSES). Examples of TSEs include Bovine Spongiform Encephalopathy (BSE), scrapic, Creutzfeldt-Jakob disease (CJD) and Gerstmann-Straussler-Sheinker syndrome (GSS). The concentration of this protein in body fluid or tissue samples may be measured by an assay of the present invention, in which a PrP epitope is captured by an antibody, which is then detected. The presence of PrP indicates BSE. PrP epitopes (AAB07320-B07326) are derived from the protease resistant core of PrP that is occluded when the PrP is in an aggregated state.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Page 42-43; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Novel immunoassay for prion protein, used transmissible spongiform encephalopathies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      WPI; 2000-387880/33.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cattle prion protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                          Cattle; prion protein; transmissible spongiform encephalopathy; bovine spongiform encephalopathy; TSE diagnosis; PrP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                17-OCT-2000
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB07328;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAB07328 standard; protein; 217 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                      Key
                                                                                                                                                                                                                                                                                                                                                                                                                 Bos bovis.
          New immunoassay for prion protein, used for determination of transmissible spongiform encephalopathies in mammals, comprispecific capture antibody .
                                                                                                                                                                                                                                                                                                                             Disulfide-bond
                                                                                                                                                                                                                                                                                                                                                                       Region
                                                                    WPI; 2000-399778/34
                                                                                               Hope J, Barnard GJR,
                                                                                                                                                                    17-NOV-1998;
                                                                                                                                                                                                 27-OCT-1999;
                                                                                                                                                                                                                             25-MAY-2000
                                                                                                                                                                                                                                                        WO200029849-A1
                                                                                                                                                                                                                                                                                                                 Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              129 GSDYEDRYY 137
                                                                                                                            (BBSR-)
                                                                                                                                         (WALL-) WALLAC OY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 GXDXEDRYY 9
                                                                                                                            BBSRC OFFICE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             217 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (first entry)
                                                                                                                                                                       98FI-0002480
                                                                                                                                                                                                  99WO-FI00896.
                                                                                                                                                                                                                                                                                                                 166..201
217
                                                                                                                                                                                                                                                                                                                                         /note= "Repeat region consisting of tandem repeats
of repeat unit: PHGGGWGQ (AAB07319)"
                                                                                                                                                                                                                                                                                                                                                                     Location/Qualifiers 37..79
                                                                                                                                                                                                                                                                                    /note= "C-terminal phospho-inositol glycolipid
membrane anchor (-GPI)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     93.2%;
                                                                                                  Birkett CR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 41; DB Pred. No. 2.6; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            for the determination of in bovines -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 217; .
                             comprises
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RESULT 14
ANDO 2561
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AC ANDO 250
A
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This sequence represents the bovine prion protein (PPP). The protein is targeted by the antibody of the invention, which is a monoclonal antibody or fragment capable of specifically binding to native and denatured normal (PPPC) and disease-specific prion protein (PPSC) in an antigen-antibody complex. The antibodies that immunoreact with disease-specific prion proteins are used in test kits for the diagnosis of prion diseases and to detect disease-specific PPP in biological material by treatment of a probe of the material with proteinase K and then with the monoclonal antibody. The monoclonal antibodies are used for
                                                                                                                                                                                                                                                                                                                                                 Disclosure; Page 20-21; 35pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                        e.g. bovine, spongiform encephalopathy, scrapic and Creutzfeldt-Jakob disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    New monoclonal antibodies specifically bind to disease-specific prion proteins - used to diagnose, prevent and treat prion diseases
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prion protein; PrP; cow; disease-specific prion protein; scraple; Kuru; prion disease detection; bovine spongiform encephalopathy; therapy; Creutzfeldt-Jakob disease; Gerstmann-Straussler-Scheinker syndrome;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The present sequence is the cattle prion protein (PrP) sequence. Conversion of the normal cellular form of prP into an aggregated, insoluble isoform is implicated in the pathogenesis of Transmissible Spongiform Encephalopathies (TSEs). Examples of TSEs include Bovine Spongiform Encephalopathy (BSE), Scraple, Creutzfeldt-Jakob disease (CJD) and Gerstmann-Straussler-Sheinker syndrome (GSS). The concentration of this protein in body fluid or tissue samples may be measured by an assay of the present invention, in which a PrP epitope is captured by an antibody, which is then detected The presence of PrP indicates TSE. PrP epitopes (AAB07320-B07326) are derived from the protease resistant core of PrP that is occluded when the PrP is in an aggregated state.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1998-449112/39.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Korth
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (ERZI-) ERZIEHUNGSDIREKTION CANTON ZURICH.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      21-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                21-FEB-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EP861900-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  13-NOV-1998
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bovine prion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 AAW70261 standard; Protein; 219 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  AAW70261;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Disclosure;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       129 GSDYEDRYY 137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Local Similarity
les 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAV33005.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Moser M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             217 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Page 42-43; 50pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               93.28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stierli B, Streit P;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 21; Length 217;
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130 GSDYEDRYY 138

1 GXDXEDRYY 9

Matches

POCGI

Similarity 7; Conserv

Conservative

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Mismatches

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                  Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
                                                                   This invention describes a synthetic polypeptide comprising at least one defined PrP (prion protein) sequence or sequences derived therefrom that are recognised by a disease specific isoform of PrP, e.g. PrP(Sc), binding substances. The new prion protein polypeptides are useful in vaccines and pharmaceuticals for treatment of, and as diagnostic agents for diagnosts of Scrapte, BSE, Kuru and Creutzfeldt-Jacob disease. The polypeptides are also useful in pharmaceutical or chemical libraries for detection of PrP(Sc)-specific agents.
                                                Sequence
                                                                                                                                                                                                                                                      New polypeptides comprising prion protein sequences - useful for diagnosis or treatment of prion diseases e.g. Scrapie, BSE and
                                                                                                                                                                                                               Claim 13; Page 6-7; 12pp; German.
                                                                                                                                                                                                                                            Creutzfeldt-Jacob disease
                                                                                                                                                                                                                                                                                                       WPI; 1999-205964/18.
                                                                                                                                                                                                                                                                                                                                    Korth C,
                                                                                                                                                                                                                                                                                                                                                                  (PRIO-) PRIONICS AG.
                                                                                                                                                                                                                                                                                                                                                                                                20-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                           20-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DE19741607-A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   the prevention and treatment of prion diseases and to clear biological material from prions. The antibodies are used to diagnose, treat and prevent e.g. bovine spongiform encephalopathy, scrapie in sheep and Creutzfeldt-Jakob disease, Gerstmann-Straussler-Scheinker syndrome, Fatal Familial Insomnia and Kuru in humans. The diagnostic method allows mass screening of infected cattle tissue at a subclinical stage and reduces possible human health risks.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                detection.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Prion protein; Prp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bovine rbPrP protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     17-JUN-1999
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           treatment; diagnosis;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         130 GSDYEDRYY 138
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nes 7; Conserv
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                                                                                                                                                                                                                                                                                                                                   Moser M,
                                               219 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          219 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                             97DE-1041607
                                                                                                                                                                                                                                                                                                                                                                                                                         97DE-1041607.
                                                                                                                                                                                                                                                                                                                                   Oesch B;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     rbPrP; disease specific isoform; PrP(Sc); vaccine;
s; Scraple; BSE; Kuru; Creutzfeldt-Jacob disease;
   93.28;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93.28;
Score 41; DB 20;
Pred. No. 2.6;
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Pred. No. 2.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           19;
               Length 219;
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Search completed: March 24, 2003, 17:19:40 Job time : 26.6875 secs



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Maximum DB
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Pred. No. 18 the number of results predicted by chance to have a
score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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44444444444
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Match
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A;Cross-references: GB:M13899; NID:g190467; PIDN:AAA60182.1; R;Puckett, C.; Concannon, P.; Casey, C.; Hood, L. Am. J. Hum. Genet. 49, 320-329, 1991 A; Residues: 1-253 < KRE> A;Title: Molecular cloning of a human prion protein cDNA. A;Reference number: A24173; MUID:86300093; PMID:3755672 A; Accession: A24173 C;Species: Homo sapiens (man) C;Date: 25-Oct-1987 #sequence_revision 12-Apr-1996 #text_change 16-Jun-2000 C;Accession: A24173; A40372; A05017; S14078; I54322; I68597; I58135; I59184; I79633; R;Kretzschmar, H.A.; Stowring, L.E.; Westaway, D.; Stubblebine, W.H.; Prusiner, S.B. major prion protein precursor - human N; Alternate names: 11K amyloid protein; 27-30K sialoglycoprotein; PrP 27-30; PrP 33-3 A;Molecule type: protein
A;Residues: 58-72,'x',74-76,'xx',79,'xxx',83-86;111-128,'V',130-150 <TAG>
A;Residues: 58-72,'x',74-76,'xx',79,'xxx',83-86;111-128,'V',130-150 <TAG>
R;Diedrich, J.F.; Knopman, D.S.; List, J.F.; Olson, K.; Frey, W.H.
Hum. Mol. Genet. 1, 443-444, 1992
A;Title: Deletion in the prion protein gene in a demented patient.
A;Reference number: 154322; MUID:93250789; PMID:1363802
A;Accession: 154322 A; Molecule type: mRNA A; Residues: 8-117,119-253 < LIA> A; Residues: B-117,119-253 < LIA> A; Cross-references: GB:D00015; NID:g220015; PIDN:BAA00011.1; PID:g220016; GB:M13667; R; Tagilavini, F.; Prelli, F.; Ghiso, J.; Bugiani, O.; Serban, D.; Prusiner, S.B.; Far EMBO J. 10, 513-519, 1991 A;Cross-references: GB:X83416; NID:g747846; PIDN:CAA58442.1; PID:g747847 A;Note: the deletion may be a polymorphism; the alternative deletion of R;Liao, Y.C.J.; Lebo, R.V.; Clawson, G.A.; Smuckler, E.A. Science 233, 364-367, 1986 A;Title: Genomic structure of the human prion protein gene A;Reference number: A40372; MUID:91328137; PMID:1678248 A;Accession: A40372 A;Cross-references: GB:M81929; NID:g190517; PIDN:AAB59442.1; PID:g190518
A;Accession: I68597
A;Status: translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 8-240 <RE3> A;Title: Amyloid protein of Gerstmann-Straeussler-Scheinker disease (Indiana kindred) A;Reference number: S14078; MUID:91160504; PMID:1672107 A;Accession: S14078 A; Molecule type: DNA A; Residues: 1-80,89-253 < PUC> A; Status: not compared with conceptual translation A; Molecule type: DNA A; Residues: 9-83,92-240 <RES> A; Status: preliminary; translated from GB/EMBL/DDBJ A; Accession: A05017 A;Reference number: A05017; MUID:86261778; PMID:3014653 1986 deletion of 82-89 could PID:g190468

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A; Gene: GDB:PRNP; CJD; PRIP
A; Gene: GDB:PRNP; CJD; PRIP
A; Gene: GDB:PRNP; GDB:120720; OMIM:176640; OMIM:137440
A; Map position: 20pter-20p12
A; Introns: #status absent
A; Note: one intron occurs before the initiator codon
A; Note: one intron occurs before with Creutzfeld-Jakob disease (CJD), Gerstmann-Strausler
C; Superfamily: major prion protein
C; Superfamily: major prion protein
C; Reywords: amylold; blocked carboxyl end; brain; glycoprotein; lipoprotein; phosphatidy
F; 1-22/Domain: signal sequence #status predicted <SIG>
F; 23-20/Product: major prion protein #status predicted <MAT>
F; 54-92/Region: #:residue repeats (P-H-G-G-W-G-Q)
F; 112-134/Domain: transmembrane #status predicted <TMI>
F; 131-253/Domain: carboxyl-terminal propeptide #status predicted <CTP>
F; 179-214/Disulfide bonds: #status predicted
F; 181,197/Binding site: carbohydrate (Asn) (covalent) #status predicted
F; 230/Modified site: GPI-anchor ethanolamine amidated carboxyl end (Ser) (in mature form
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A;Title: Infectious amyloid precursor gene sequences in primates used for experimental A;Reference number: I36907; MUID:95083661; PMID:7991800
A;Accession: I37032
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A;Residues: 1-253 <RES>
A;Cross-references: EMBL:U15166; NID:g563208; PIDN:AAA68633.1; PID:g563209
C;Superfamily: major prion protein
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C;Species: Gorilla gorilla (gorilla)
C;Date: 31-May-1996 #sequence_revision 31-May-1996 #text_change 13-Aug-1999
C;Accession: I37032
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R;Brown, P.; Goldfarb, L.G.; McCombie, W.R.; Nieto, A.; Squillacote, D.; Sheremata, W.;
Neurology 42, 422-427, 1992
A;Title: Atypical Creutzfeldt-Jakob disease in an American family with an insert mutatic
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C;Genetics:
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A; Residues: 60-67 <GOL>
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A; Accession: I59184
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A;Accession: I58135
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RESULT 3

major prion protein - common gibbon

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A;Cross references: EMBL:U08308; NID:g474374; PIDN:AAC50096.1; PID:g474375 A;Note: the source was designated as Symphalangus syndactylus A;Note: the nucleotide sequence was submitted to the EMBL Data Library, Ap:C;Superfamily: major prion protein
                                                                                                                                                                                                                                                                                                                                                                                                    R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B. J. Mol. Biol. 245, 362-374, 1995
A;Title: Prion protein gene variation among primates.
A;Reference number: S53614; MUID:95139066; PMID:7837269
A;Accession: S53635
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      prion protein - siamang
C:Species: Hylobates syndactylus (siamang)
C;Date: 15-Jul-1995 #sequence_revision 19-Apr-1996 #text_change 13-Aug-1999
C;Accession: S53635
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A; Residues: 1-253 <SCH>
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C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane
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C:Species: Pan troglodytes (chimpanzee)
C:Date: 31-May-1996 #sequence_revision 31-May-1996 #text_change 13-Aug-1999
C:Accession: I61847; S71060; S53615
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R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.
J. Mol. Biol. 245, 362-374, 1995
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A; Residues: 1-253 <SCW>
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A; Residues: 1-210, 'R', 212-253 <SCH>
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A;Molecule type: DNA
A;Residues: 1-253 <RES>
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Pred. No. 0.42;
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Pred. No. 0.42;
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C;Species: Hylobates lar (common gibbon, white-handed gibbon)
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 13-Aug-1999
C;Accession: $53617; $71050
R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.
J;Mol. Biol. 245, 362-374, 1995
A;Title: Prion protein gene variation among primates.
A;Reference number: $53614; MUID:95139066; PMID:7837269
A;Accession: $53617
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A;Residues: 1-210,'E',212-253 <SCW>
A;Residues: 1-210,'E',212-253 <SCW>
A;Cross-references: EMBL:U0829; NID:g474356; PIDN:AAC50088.1; PID:g474357
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
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A; Accession: S71050
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A;Residues: 1-210,'E',212-253 <SCW>
A;Residues: 1-210,'E',212-253 <SCW>
A;Cross-references: EMBL:U08300; NID:g474358; PIDN:AAC50089.1; PID:g474359
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane
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                                                                  C;Species: Tragelaphus strepsiceros (greater kudu)
C;Date: 06-Jan-1995 #sequence_revision 06-Jan-1995
C;Accession: S37137
                                                                                                                                                                                                                                                                                                          밁
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Reference number: S71041
A; Accession: S71049
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          submitted to the EMBL Data Library, April 1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Residues: 1-253 <SCH>
A;Cross-references: EMBL:U08300
n;martin, T.C.; Hughes, S.L.; Hughes, K.J.; Dawson, M. submitted to the EMBL Data Library, August 1993
                                                                                                                                                         prion protein - greater kudu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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les 7; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GXDXEDRYY 9
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77.88;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Mismatches
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Pred. No. 0.42;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 41; DB 2; Length 253; Pred. No. 0.42;
                                                                                                                                                                                                                                                                                                                                                                                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           07-Feb-1997 #text_change 13-Aug-1999
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                                                                                                         #text_change 13-Aug-1999
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major prion protein 1 precursor - bovine
N;Alternate names: prion protein, long variant; Prp protein
C;Species: Bos primigenius taurus (cattle)
C;Date: 09-Sep-1994 #sequence_revision 09-Sep-1994 #text_change 02-Mar-2001
C;Accession: A54330; JT0953; JT0952; A48551; S07347; I46931
C;Accession: A54330; JT0953; Martin, T.; Dawson, M.; Hope, J.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Molecule type: DNA
A;Residues: 1-264 <MAR>
A;Cross-references: EMBL:X74771; NID:g398937; PIDN:CAA52781.1; PID:g398938
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Reference number: S37137
A; Accession: S37137
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Note: sequence extracted from NCBI backbone (NCBIN:121620, NCBIP:121621)
R; Hope, J.; Reekle, L.J.D.; Hunter, N.; Multhaup, G.; Beyreuther, K.; White, H.; Scot Nature 336, 390-392, 1988
A; Title: Fibrils from brains of cows with new cattle disease contain scraple-associat A; Reference number: S07347; MUID:89057122; PMID:2904126
A; Accession: S07347
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A;Gene: prP
C;Superfamily: major prion protein
C;Keywords: glycoprotein; phosphatidylinositol linkage;
F:1-24/Domain: signal sequence #status predicted <SIG>
                                                                                                                                                                                                                                                                                                                                                           A;Molecule type: protein
A;Molecule type: protein
A;Residues: 25-36 <HOP>
A;Residues: 25-36 <HOP>
R;Prusiner, S.B.; Fuzi, M.; Scott, M.; Serban, D.; Serban, H.; Taraboulos, A.; Gabrie
R;Prusiner, Dis. 167, 602-613, 1993
J. Infect. Dis. 167, 602-613, 1993
A;Title: Immunologic and molecular biologic studies of prion proteins in bovine spong
A;Reference number: 146931; MUID:93179783; PMID:8440932
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Title: Comparative sequence analysis and expression of bovine PrP A;Reference number: A48551; MUID:93118243; PMID:1362024 A;Accession: A48551
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A; Molecule type: mRNA
A; Residues: 1-217, 'K', 219-264 < YO3>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Yoshimoto, J.; Iinuma, T.; Ishiguro, N.; Horiuchi, M.; Imamura, M.; Shinagawa, Virus Genes 6, 343-356, 1992
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Molecule type: DNA
A; Residues: 1-217, 'K
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Residues: 1-264 <YOS>
A;Residues: 1-264 <YOS>
A;Cross-references: GB:D10613; NID:g217595; PIDN:BAA01468.1; PID:g217596
A;Accession: JT0952
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R; YOShimoto, J.; Ilnuma, T.; Ishi
submitted to JIPID, November 1991
A; Reference number: JT0952
A; Accession: JT0953
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-264 <GOL>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  R; Goldmann, W.; Hunter, N.;
J. Gen. Virol. 72, 201-204,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A; Molecule type: DNA
                                                                                                                                                                                                                                        A; Molecule type: mRNA
A; Residues: 1-264 < PRU>
                                                                                                                                                                                                                                                                                                 A;Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                   A; Accession: I46931
                                                                                                                                                                                                          A;Cross-references: GB:S55629; NID:g266111; PIDN:AAB25514.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
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the bovine PrP gene have five or six copies of a short, G
MUID:91116314; PMID:1671225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ishiguro,
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                                                                                polymorphism; tandem repeat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Imamura, M.;
                                                                                                                                                                                                             PID:g266112
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           gene in mouse L-9
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A;Molecule type: DNA
A;Residues: 1-202, E',204-239 <SCW>
A;Residues: 1-202, E',204-239 <SCW>
A;Cross-rences: EMBL:U08293; NID:g474344; PIDN:AAC50082.1; PID:g474345
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Cross-references: EMBL:U08309; NID:9474376; PIDN:AAC50097.1; PID:9474377
R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.

J. Mol. Blol. 245, 362-374, 1995
A;Title: Prion protein gene variation among primates.
A;Reference number: S53614; MUID:95139066; PMID:7837269
A;Accession: S53630
A;Status: nucleic acid sequence not shown
A;Molecule type: DNA
A;Residues: 1-194,'R',196-231 <SCW>
A;Cross-references: EMBL:U08309
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
                                                                                                                                                                                                                                                                                                                                                                             major prion protein - douroucouli (fragment)
C;Species: Aotus trivirgatus (douroucouli, night monkey, owl monkey)
C;Date: 28-Oct.1996 #sequence_revision 07-Feb-1997 #text_change 13-Aug-1999
C;Accession: S53633; S71042
R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.
J. Mol. Biol. 245, 362-374, 1995
A;Title: Prion protein gene variation among primates.
A;Reference number: S53614; MUID:95139066; PMID:7837269
A;Accession: S5363
                                                                                                                                                                     submitted to the EMBL Data Library, April 1994 A; Reference number: S71041 A; Accession: S71042
                                                                                                                                                                                                                                                                                                            A;Status: nucleic acid sequence not shown A;Molecule type: DNA A;Residues: 1-239 <SCH>
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A; Residues: 1-232 <SCH>
A; Cross-references: EMB
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  mmjor prion protein - black-handed spider monkey (fragment)
C;Specles: Ateles geoffroy1 (black-handed spider monkey)
C;Date: 27-Oct-1996 #sequence_revision 07-Feb-1997 #text_change 13-Aug-1999
C;Accession: S71041; S53630
C;Accession: H.M. SS3630
Submitted to the EMBL Data Library, April 1994
A;Reference number: S71041
                                                                                                                                                                                                                                                                                  A;Cross-references: EMBL:U08293
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F;192,208/Blnding site: carbohydrate (Asn) (covalent) #status predicted
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F;60-99/Region: 8-residue repeats (W-G-Q-P-H-G-G-G)
F;190-225/Disulfide bonds: #status predicted
F:100-205/Disulfide bonds: #status predicted
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Best Local (
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GXDXEDRYY 9
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77.8%;
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Pred. No. 0.62;
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Pred. No. 0
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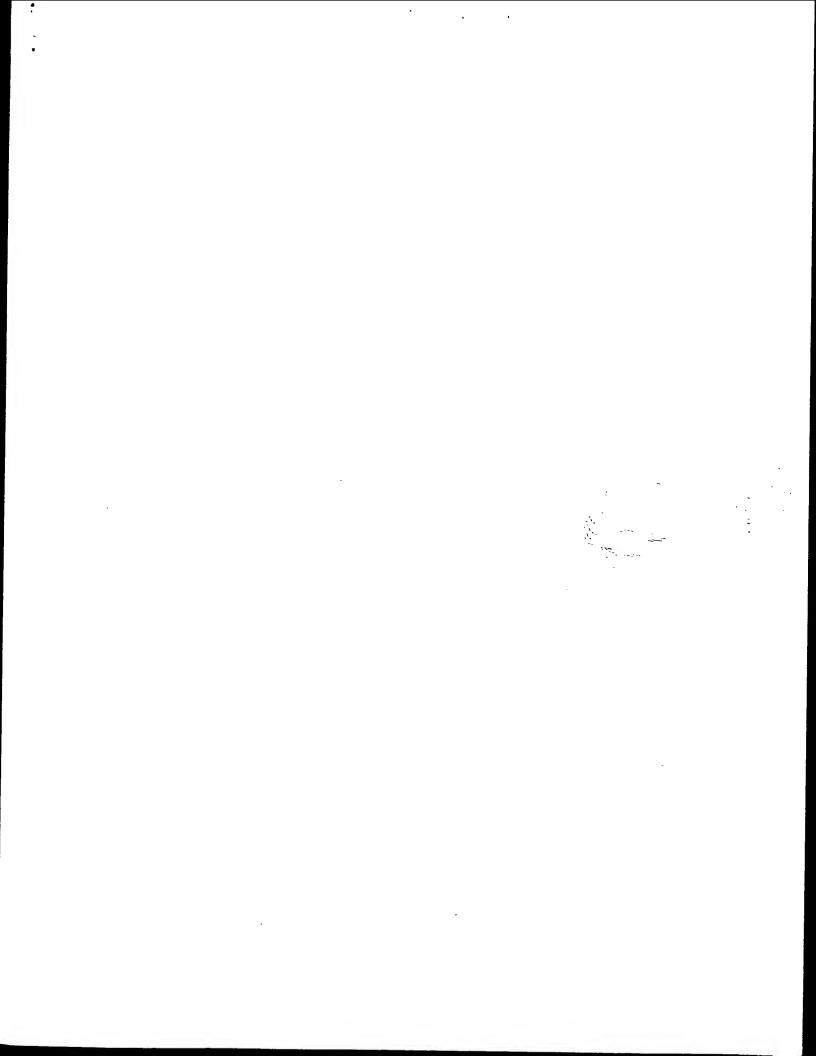
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A Status: nucleic acid sequence not shown A; Molecule type: DNA A; Molecule type: DNA A; Cross-references: EMBL: U08312 C; Superfamily: major prion protein C; Superfamily: major prion pr
                                                                                                                                                C; Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                 A;Reference number: S53614; MUID:95139066; PMID:7837269
A;Accession: S53632
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: EMBL:U08312; NID:g475585; PIDN:AAC50100.1; PID:g475586 R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B. J. Mol. Biol. 245, 362-374, 1995
A;Title: Prion protein gene variation among primates.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              R;Schatzl, H.M.
submitted to the EMBL Data Library, April 1994
A;Reference number: S71041
A;Accession: S71048
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C:Species: Callicebus moloch
C:Date: 27-Oct-1996 #sequence_revision 07-Feb-1997 #text_change 13-Aug-1999
C:Accession: S71048; S53632
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C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A; Molecule type: DNA
A; Residues: 1-241 <SCH>
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A;Reference number: S53614; MUID:95139066; PMID:7837269 A;Accession: S53621
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A; Residues: 1-241 <SCH>
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A; Accession: S71056
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C;Species: Papio sphinx, Mandrillus sphinx (mandrill)
C;Date: 27-Oct-1996 #sequence_revision 14-Feb-1997 #t.
C;Accession: S71056; S53621
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                                                                                        Query Match
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Matches 7; Conserv
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                                                Local
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                Conservative
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                                        90.9%;
77.8%;
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77.8%;
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                                        Score 40; DB 2
Pred. No. 0.64;
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Pred. No. 0.64;
        Mismatches
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                                                                                DB 2;
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0.64;
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                                                                            Length 241;
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Gaps
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major prion protein - green monkey
C; Species: Cercopithecus aethiops (green monkey, grivet)
C; Apate: 28-Oct-1996 *sequence_revision 07-Feb-1997 *text_change 13-Aug-1999
C; Accession: S53627; S71043
R; Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.
J. Mol. Biol. 245, 362-374, 1995
J. Mol. Biol. 245, 362-374, 1995
A; Areference number: S53614; MUID:95139066; PMID:7837269
A; Accession: S53627
A; Status: nucleic acid sequence not shown
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S71045
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A; Residues: 1-10, 'v',12-202, 'E',204-245 <SCW>
A; Residues: 1-10, 'v',12-202, 'E',204-245 <SCW>
A; Cross-references: EMBL:U08291; NID:g474340; PIDN:AAC50080.1; PID:g474341
C; Superfamily: major prion protein
C; Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
                                                                                                                                                                                                                                                                                                                                                                    A:Cross-references: EMBL:U08292; NID:9474342; PIDN:AAC50081.1; PID:9474343 R:Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B. J. Mol. Biol. 245, 362-374, 1995 A:Title: Prion protein gene variation among primates. A:Reference number: S53614; MUID:95139066; PMID:7837269 A:Accession: S53628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         major prion protein - Cercopithecus dianae
C:Species: Cercopithecus dianae
C:Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 13-Aug-1999
C:Accession: S71045; S53628
                                                                                                                                                                                                      A;Status: nucleic acid sequence not shown
A;Molecule type: DNA
A;Residues: 8-10,'L',12-202,'R',204-239 <SCW>
A;Cross-references: EMBL:U08292
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
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A; Residues: 1-245 <SCH>
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134 GNDYEDRYY 142
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Local Similarity 77.8%;
                                                                                              Local Similarity es 7; Conserv
                                                   1 GXDXEDRYY 9
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                                                                                                    Conservative
                                                                                                                              90.9%; Score 40; DB 2; Length 245; 77.8%; Pred. No. 0.65;
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                                                                                                             2; Indels
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R;Cervenakova, L.; Brown, P.; Goldfarb, L.G.; Nagle, J.; Pettrone, K.; Rubenstein, R. Proc. Natl. Acad. Sci. U.S.A. 91, 12159-12162, 1994
A;Title: Infectious amyloid precursor gene sequences in primates used for experimenta A;Reference number: 136907; MUID:95083661; PMID:7991600
A;Accession: I61848
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         major prion protein precursor - common squirrel monkey
C;Species: Saimiri sciureus (common squirrel monkey)
C;Date: 31-May-1996 #sequence_revision 31-May-1996 #text_change 13-Aug-1999
C;Accession: I61848
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A;Residues: 1-252 <RES>
A;Cross-references: EMBL:U15165; NID:g595852; PIDN:AAA68636.1; PID:g595853
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141 GNDYEDRYY 149
                                                                                                            MAICH 90.9%; Score 40; DB 2; Length 252; Local Similarity 77.8%; Pred. No. 0.68; es 7; Conservative 0. Mismath.
                                                            1 GXDXEDRYY 9
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Search completed: March 24, 2003, 17:24:13 Job time: 12.25 secs



GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Scoring table:
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                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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RA Beasley O.P., Bird C.P., Blakey S.E., Bridgeman A.M., Brown A.J.,
RA Buck D., Burrill W.D., Butler A.P., Carder C., Carter N.P.,
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RA Culson A., Coville G.J., Deadman R., Dhami P.D., Dunn M.,
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RA Hammond S., Harley J.L., Heath P.D., HO S., Holden J.L., Howden P.J.,
RA Huckle E., Hunt A.R., Hunt S.E., Jekosch K., Johnson C.M., Johnson D.,
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Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C) (ASCR)
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"Molecular cloning of a human prion protein cDNA.";
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"Amyloid protein of Gerstmann-Straussler-Scheinker disease (Indiana kindred) is an 11 kd fragment of prion protein with an N-terminister of prion protein with a N-terminister of prion prion prion prion prion prion prion prion p
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MEDLINE-92195483; PubMed-1347910;
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syndrome.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-93372867; PubMed-8364585; Palmer M.S., Collinge J.; Mutations and polymorphisms in the Hum. Mutat. 2:168-173(1993).
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Zahn R., Liu A., Luhrs T., Riek R., von Schroetter C.,

Lopez Garcia F., Billeter M., Calzolai L., Wider G., I

"NMR solution structure of the human prion protein.";

"NMR solution structure of the human prion protein.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                         Nature 338:342-345(1989).
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MEDLINE-20359708; PubMed-10900000;
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Science 233:364-367(1986).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE OF 8-253 FROM N.A.
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                                                                                                                                                                                                                                                                               but not the sole mutation related to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   at codon 58.";
10:513-519(1991).
42:669-670(1992).
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Allcatelli C.T., Masullo C., Macchi G., Alema
Petraroli R., Silvestrini M.C., Brunori M.;
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Tateishi J., Otomo E., Hayakawa M., Tanaka J., Matsushita M.,
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MEDLINE-91124933; PubMed-1671440;
                                                      MEDLINE-94316708; PubMed-7913755;
Gabizon R., Rosenman H., Meiner Z., Kahana I., Kahana
Ott J., Prusiner S.B.;
                                                                                                                                                    VARIANT CJD LYS-200.

MEDLING-94142912; PubMed=7906019;

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mutation of the prion protein gene.";
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MEDLINE-93213314; PubMed-8461023;
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MEDLINE-95213742; PubMed-7699395;
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Jakob disease with codon 200 po
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ema G., Galgani S., Xi Y.G.,
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RESULT 3
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Young K., Jones C.K., Piccardo P., Lazzarini A., Golbe L.I.,
Zimmerman T.R., Dickson D.W., McLachlan D.C., St George-Hyslop P.H.,
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Proc. Natl. Acad. Sci. U.S.A. 91:12159-12162(1994).
-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBJUIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Neurology 45:1127-1134(1995).
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Schattl H.M., Dacosta M., Taylor L.,
"Prion protein gene variation among
J. Mol. Biol. 245:362-374(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Hylobates lar (Common gibbon), and Hylobates syndactylus (Siamang) (Symphalangus syndactylus). Hylobates syndactylus (Siamang) (Symphalangus syndactylus). Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Pan. NCBI_TaxID-9598, 9580, 9590;
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01-FEB-1995 (Rel. 31, Last sequence update)
30-MAY-2000 (Rel. 39, Last annotation update)
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                                                                                                                                                                                                                SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                   SIMILARITY: BELONGS TO THE PRION FAMILY.
U08296; AAC50085.1; -. U08299; AAC50088.1; -.
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EMBL; U08308; EMBL; U15039; HSSP; P04156;

AAC50096.1; AAA68632.1;

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                                                                                                                                                                                                                                       Martin T.C., Hughes S.L., Hughes K.J., Dawson M.;
Submitted (AUG-1993) to the EMBL/GenBank/DBJ databases.
-I- FUNCTION: THE FUNCTION OF PRP IS NOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                        Tragelaphus strepsiceros (Greater kudu).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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01-FEB-1995 (Rel. 31, Last sequence update)
01-FEB-1995 (Rel. 31, Last annotation update)
Major prion protein 2 precursor (PrP) (Major scrapie-associated fibril
                                                                                                                                                                                                                                                                                                                        TISSUE-Brain;
                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
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REPEAT
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P40243;
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                                                                                                                              SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH DEGENERATIVE NEUROLOGICAL DISEASES SUCH AS KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                      SIMILARITY: BELONGS TO THE PRION FAMILY.
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7; Conser
email to license@isb-sib.ch).
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27633 MW;
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Pred. No. 0.2;
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BY SIMILARITY.
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N-LINKED (GLCNAC. . .) (POTENTIAL).
5 X 8 AA TANDEM REPEATS OF P-H-G-G-
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REMOVED IN MATURE FO
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Best Local
                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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01-OCT-1996
01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                           Martin T., Hughes S., Hughes K., Dawson M.;
"Direct sequencing of PCR amplified pig PrP genes.";
Blochim. Blophys. Acta 1270:211-214(1995).
-1- FUNCTION: THE FUNCTION OF PRP IS OUT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-1- SUBJUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIO_PIG P49927;
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CARBOHYD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Cetartiodactyla; Suina; Suidae;
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PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
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Les 7; Conserv
                                                                                                                                                                                                                                             SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIN INFECTED WITH DEGENERATIVE NEUROLOGICAL DISEASES SUCH AS KURU, CREUTZFELDT-LAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
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34, Last annotation update)
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Pred. No. 0
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BY SIMILARITY.
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EMBL; L07623; AAA92862.1; -. .

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoldea
Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOY-1991 (Rel. 20, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
Major prion protein 1 precursor (PrP) (Major scrapie-associated fibril
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       BOVIN
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                                    SEQUENCE FROM N.A.

MEDLINE-93179783; PubMed-8440932;

MEDLINE-93179783; PubMed-8440932;

Prusiner S.B., Fuzi M., Scott M., Serban D., Serban H.,

Prusiner S.B., Fuzi M., Scott M., Serban D., Serban H.,

Taraboulos A., Gabriel J.M., Wells G.A., Wilesmith J.W., Bradl

Tamunologic and molecular biologic studies of prion proteins
bovine spongiform encephalopathy.";

J. Infect. Dis. 167:602-613(1993).
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Goldmann W., Hunter N., Martin T., "Different forms of the bovine PrP short, G-C-rich element within the
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REPEAT
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                                                                                                                                                                                                                              "Comparative sequence analysis and expression mouse L-929 cells.";
Virus Genes 6:343-356(1992).
                                                                                                                                                                                                                                                                                              Shinagawa M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STRAIN-Holstein-Friesian;
MEDLINE-91116314; PubMed-1671225;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                          TISSUE-Brain;
                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
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SEQUENCE FROM N.A.
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7; Conser
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (Rel.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3
                                                                                                                                                                                                                                                                                                                                                                                                                                          72:201-204(1991).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRP;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    . 10, Created)
. 20, Last sequence. 41, Last annotation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRION.
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70
78
86
85
95
27727
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93.2%;
                                                                                                                                                                                                                                                                                                                         Ŧ.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   .88;
                                                                                                                                                                                                                                                                                                                         Ishiguro N., Horiuchi M., Imamura M.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ¥.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   برب
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 41; DB:
Pred. No. 0.2;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             GPI-anchor; Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MAJOR PRION PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3A87104B234C55DD CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AA TANDEM REPEATS OF P-H-G-G-G-W-G-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  protein-coding exon.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Dawson M., Hope J.;
gene have five or six copies of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    264 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2;
                                                                                                                                                                                                                                                                               of bovine PrP gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 257;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Signal
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                                                                                                                Bradley
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REAL BREEF B
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                                          Pfam; PF00377; prion; 1.
PRINTS; PR00341; PRION.
SMART; SM00086; PAC; 1.
SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                    PDB; 1DX0; 20-JUL-00.
PDB; 1DX1; 20-JUL-00.
InterPro; IPR001610; PAC.
InterPro; IPR000817; Prion.
Pfam; PF00377; Prion; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE OF 25-36.

MEDLINE-89057122; PubMed-2904126;

Hope J., Reekie L.J.D., Hunter N., Multhaup G., Beyreuther K.,

White H., Scott A.C., Stack M.J., Dawson M., Wells G.A.;

White H., Scott B.G., Stack M.J., Dawson M., Wells G.A.;

White H., Scott B.G., Stack M.J., Dawson M., Wells G.A.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation—the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Complete genomic sequence of the bovine polymorphism in its promoter region."; Anim. Genet. 32:231-232(2001).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hills D., Comincini S., Schlaepfer J., Williams J.L.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-20359707; PubMed-10899999;
Lopez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Wuethrich K.;
Impez Garcia F., Zahn R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Riek R., Wuethrich K.;
Impez Garcia F., Zahn R., Wuethrich K.;
Impez Garcia F., Wuethrich F., Wuethrich K.;
Impez Garcia F., Wuethrich K.;
Impez Garcia 
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STRAIN-Holstein-Friesian; TISSUE-Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     associated protein.";
Nature 336:390-392(1988).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Tanaka M., Inoue S., Ikeda T., Horiuchi M., Ishiguro N., Shinagawa M.; Submitted (JAN-1994) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE OF 1-15 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MEDLINE-21422903; PubMed-11531705;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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Prion; Brain; Glycoprotein; GPI-anchor; Repeat; Signal;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DISEASE: FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH DEGENERATIVE NEUROLOGICAL DISEASES SUCH AS KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             20-JUL-00.
20-JUL-00.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Dolf G.,
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          3D-structure
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local
                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
01-NOV-1997 (Rel. 35, Last annotation update)
Major prion protein 1 precursor (PrP) (Major scrapie-associated f. Protein 1).
Tragelaphus strepsiceros (Greater kudu).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoldea;
Bovidae; Bovinae; Tragelaphus.
              Pram; PF00377; prion; 1. PRINTS; PR00341; PRION. SMART; SM00157; PRP; 1.
                                         InterPro; IPR000817; Prion. Pfam; PF00377; prion; 1.
                                                                                    PIR;
                                                                                                                                                                                                                                                                                                                                       MARTIN T.C., Hughes S.L., Hughes K.J., Dawson M.;
Submitted (AUG-1993) to the EMEL/GenBank/DBJ databases.
-I- FUNCTION: THE FUNCTION OF PRP IS NOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
    PROSITE;
                                                                      HSSP; P10279; 1DWY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CONFLICT
                                                                                               EMBL;
                                                                                                                                                                                                                                                                                                                                                                                                  TISSUE-Brain;
Martin T.C.,
                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                        NCBI_TaxID-9946;
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P40242;
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REPEAT
REPEAT
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REPEAT
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CHAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         1 GXDXEDRYY 9
                                                                                                                                                                                                                                         SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH DEGENERATIVE NEUROLOGICAL DISEASES SUCH AS KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                             s37137; s37137.
                                                                                                                                                                                                                                  SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
7; Conserv
  PS00291;
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PRION_1; 1.
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Pred. No. 0.21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MISSING (IN A SECOND ALLELE).
E -> K (IN REF. 2 AND 4).
D6D214038316A231 CRC64;
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N-LINKED (GLCNAC. . .)
N-LINKED (GLCNAC. . .)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8
8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 tation update)
(PrP) (Major scraple-associated fibril
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     AA TANDEM REPEATS OF P-H-G-G-W-G-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      264 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 264;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PROBABLE).
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PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
Prion: Rrain: Glycoprotein; GPI-anchor; Repeat; Signal.

Pfam; PF00377; prion; 1 SMART; SM00157; PRP; 1.

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RESULT 8
PRIO_ATEGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P40246;
01-FEB-1995 (
01-FEB-1995 (
01-OCT-1996 )
                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDLINE-95139066; PubMed-7837269;
Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;
"Prion protein gene variation among primates.";
J. MOI. Biol. 245:362-374(1995).
-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REPEAT
SEQUENCE
                   EMBL; U08309; AAC50097.1; -. HSSP; P04156; 1E1G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Ateles geoffroyi (Black-handed spider monkey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Platyrrhini; Cebidae;
                                                                                                                                                                                                                                                                                                                                                                                                                        HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND IN-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Major prion
InterPro; IPR000817; Prion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID-9509;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIO_ATEGE
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REPEAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          153 GSDYEDRYY 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REPEAT
                                                                                                                                                                                                                                                                 SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                          MOI. Biol. 245:362-374(1995).
FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS
                                                                                                                                                                                                                                                    SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Similarity 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (Rel. 34, Last annotation update) protein precursor (PrP) (PrP27-30)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           (Rel. 31, Created)
(Rel. 31, Last sequence update)
(Rel. 34, Last annotation updat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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1 24
5 264
5 192
2 192
8 208
2 25
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77.8%;
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78
86
94
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POTENTIAL.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Score 41; DB 1;
Pred. No. 0.21;
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N-LINKED (GLCNAC. . .) (POTENTIAL).

N-LINKED (GLCNAC. . .) (POTENTIAL).

BY SIMILARITY.

6 X 8 AA TANDEM REPEATS OF P-H-G-G-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FEB73F41732619B1 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                X 8 AA TANDEM REPEATS OF P-H-G-G-W-G-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Euteleostom1;
Atelinae; Ateles
                                                                                                                                                                                                                                                                                                                                                                                                                        POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0;
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Best Local
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REPEAT
REPEAT
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DISULFID
                                                                                                                                                                                                                                                                                                                                              der Ruyl A.C., Dekker J.T., Goudsmit J.;
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIO_CERAT STANDARD; PRT; 238 AA 095145; Q95200; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation updat Major prion protein precursor (PrP) (PrP27-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NON_TER
SIGNAL
CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                  EMBL; U75384; AAB50623.1; -. EMBL; U75382; AAB50629.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cercopithecinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cercocebus aterrimus, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRNP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CERAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DOMAIN
                        Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
                                                                                                                        or send an email to license@isb-sib.ch).
                                                                                                                                      entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID-36222,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Macaca sylvanus (Barbary ape).
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                                                       InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      126 GNDYEDRYY 134
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 GXDXEDRYY 9
                                                                                                                                                                                                                                      SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                          SUBCELLULAR LOCATION: Attached to the membrane by a GPT anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME
                                                                                                                                                                                                                                                 (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Similarity 7; Conser
PS00291; PRION_1; PS00706; PRION_2;
                                                       IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
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>232
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59
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MAJOR PRION PROTEIN.

REMOVED IN MATURE FORM (BY SI
GPI-ANCHOR (BY SIMILARITY).

BY SIMILARITY.

N-LINKED (GLCNAC. . . .) (POTEN
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Pred. No.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           notation update) (PrP) (PrP27-30) (PrP33-35C) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0E2D75F04C05CC4A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              KED (GLCNAC. . .) (POTENTIAL).
KED (GLCNAC. . .) (POTENTIAL).
AA TANDEM REPEATS OF P-H-G-G-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
0.29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  FORM (BY SIMILARITY).
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Best Local S
Matches 7
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SEQUENCE
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DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prion; I
NON_TER
SIGNAL
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01-NOV-1997 (Rel.
01-NOV-1997 (Rel.
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REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CHAIN
                                                                                                                                                                                                                                                                                                                                                                    der Kuyl A.C., Dekker J.T., Goudsmit J.;
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBJUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Theropithecus gelada (Gelada baboon).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIO_THEGE Q95270;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CARBOHYD
                                                                                                                                                    use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Cercopithecinae; Theropithecus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Major prion PRNP OR PRP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CARBOHYD
          PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
Prion; Brain; Glycoprotein; G
                                                                                            EMBL; U75383; AAB50630.1;
HSSP; P04925; lAG2.
                                                                                                                               or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REPEAT
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                                                     Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=9565;
                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE FROM N.A.
                                                                                  InterPro; IPR000817; Prion.
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                                                                                                                                                                                                                                                     SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HEIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRAUSMISSIBLE MINK ENCEPHALOPATHY (TMC), ETC.
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          Glycoprotein; GPI-anchor; Repeat; Signal.
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Pred. No.
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N-LINKED (GLCNAC. . .) (POTENTIAL).
4 X 8 AA TANDEM REPEATS OF P-H-G-G-
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REMOVED IN MATURE FORM (BY GPI-ANCHOR (BY SIMILARITY)
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(PrP27-30)
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Best Local S
Matches 7
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MEDLINE-95139066; PubMed-7837269;

MEDLINE-95139066; PubMed-7837269;

Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;

Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;

"Prion protein gene variation among primates.";

J. Mol. Biol. 245:362-374(1995),

J. Mol. Biol. 245:362-374(1995),

J. MOI. B
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01-FEB-1995 (
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                                                                                                                               PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                             Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
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                                                                                                                                                                                                                                                                                                                                                                                        use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Platyrrhini; Cebidae; Aotinae; Aotu
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SWISS-PROT entry is copyright. It is produced through een the Swiss Institute of Bioinformatics and the Excurpean Bioinformatics Institute. There are no restably non-profit institutions as long as its content
                                                                                             Brain; Glycoprotein;
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MAJOR PRION PROTEIN.
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                                                                                         GPI-anchor; Repeat; Signal.
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Pred. No. 0.3;
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01-FEB-1995
01-NOV-1995
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-!- DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS ALE ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KULL CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                              "Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;
"Prion protein gene variation among primates.";
J. Mol. Biol. '245:362-374(1995).
-I- FÜNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 REPEAT
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                                                   PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                              Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
                                                                                                                    EMBL; U08312; AAC50100.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                                                     the European Bioinformatics Institute. There are no rest
use by non-profit institutions as long as its content
modified and this statement is not removed. Usage by ar
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Callicebus moloch (Dusky titi).
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Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).
                                                                                                                                                                         entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
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Mammalia; Eutheria; Primates; Platyrrhini; Cebidae; Callicebinae;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-95139066; PubMed-7837269;
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                                                                                                                                                                                                                                                                       SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                      "RODS"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CALMO
                                      Brain;
                                                                                                                                                            an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity 7; Conserv
                                                                                                          IPR000817; Prion.
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(Rel.
(Rel.
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161
                                    Glycoprotein;
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60
76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               31, Created)31, Last sequence update)32, Last annotation updat
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189
189
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67
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77.88;
                                   GPI-anchor; Repeat; Signal
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Pred. No.
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N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
5 x 8 AA TANDEM REPEATS OF P-H-G-G-W-G-Q.
1.
2.
3.
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                                                                                                                                                                                                              There are no restrictions
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        DB 1;
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Best Local
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01-FEB-1995
01-OCT-1996
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NON_TER
                                               Pfam; PF00377; prion; 1.
smart; sm00157; prp; 1.
pr051TE; PS00291; prion_1; 1.
pr051TE; PS00706; prion_2; 1.
                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mandrillus sphinx (Mandrill) (Papio sphinx).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REPEAT
REPEAT
REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE=95139066; PubMed=7837269;
Schatzl H.M., Dacosta M., Taylor L., Cohen F.E.,
"Prion protein gene variation among primates.";
J. Mol. Biol. 245:362-374(1995).
                         NON_TER
                                                                                                                EMBL; U08303; AAC50091.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                        or send
                                                                                                                                                                               modified and this statement is not removed.
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P40255;
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                                       Prion;
                                                                                                    InterPro; IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9561;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 135 GNDYEDRYY 143
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                                                                                                                                                                                                                                                          (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC. SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED MITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME
                                                                                                                                                                                                                                                                                                                                                                            MOL. B101. 245:362-374(1995).
FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
                                                                                                                                                                                                                                                                                                                                                                 SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                    Brain; Glycoprotein;
                                                                                                                                                     s requires a license agreement (See http://www.isb-sib.ch/announce/
an email to license@isb-sib.ch).
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7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (Rel.
(Rel.
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174
190
44
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61
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31, Last sequence update)
34, Last annotation updat
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174
190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              precursor (PrP)
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68
76
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N-LINKED (GLCNAC. ..) (POTENTIAL).

N-LINKED (GLCNAC. ..) (POTENTIAL).

5 X 8 AA TANDEM REPEATS OF P-H-G-G-W-G-Q.

1.
2.
3.
3.
4.
                                  GPI-anchor; Repeat; Signal.
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Pred. No.
BY SIMILARITY.
MAJOR PRION PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C6D2013EE7CAEC93 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mismatches
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(PrP27-30) (PrP33-35C) (Fragment).
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                                                                                                                                                                                Usage by
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 241;
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Pfam; PF00377; prion; 1 PRINTS; PR00341; PRION. SMART; SM00157; PRP; 1.

InterPro;

1925; 1AG2. IPR000817; Prion.

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Best Local
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                                                                                                                                                                                                                                                                                                                                                                                                              Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;
"Prion protein gene variation among primates.";
"J., Mol. Biol. 245:362-374(1995).
"I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
"BOOLIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CARBOHYD
DOMAIN
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                                                                                                This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REPEAT
NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CERAE
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REPEAT
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LIPID
EMBL; U08291; AAC50080.1; -.
EMBL; U08292; AAC50081.1; -.
HSSP; P04925; IAG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cercopithecus aethiops (Green monkey) (Grivet), Cercopithecus diana (Diana monkey).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Major prion PRNP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-FEB-1995
16-OCT-2001
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01-FEB-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                     or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE=95139066; PubMed=7837269;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9534, 36224;
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                                                                                                                                                                                                                                                       (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC. SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Rel. 31, Created)
(Rel. 31, Last sequence update)
(Rel. 40, Last annotation update)
(Rel. 40, Cast annotation update)
(Rel. 40, Cast annotation update)
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64
69
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60
68
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N-LINKED (GLCNAC. ..) (POTENTIAL)
X 8 AA TANDEM REPEATS OF P-H-G-G
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REMOVED IN MATURE FORM (BY SIMILARITY).
GPI-ANCHOR (BY SIMILARITY).
BY SIMILARITY.
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Best Local
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REPEAT
SEQUENCE
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DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  der Kuyl A.C., Dekker J.T., Goudsmit J.;
Submilted (NOV-1996) to the EMBL/GenBank/DBJ databases.

-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.

-I- SUBBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REPEAT
REPEAT
                                          InterPro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
SMART; SM00157; PRP; 1.
                                                                                                                                       EMBL; U75386; AAB50625.1; -. EMBL; U75387; AAB50626.1; -.
                                                                                                                                                                                                                                                           the European Bioinformatics Institute. The use by non-profit institutions as long modified and this statement is not removed.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   <del>-</del> -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                 HSSP; P04925;
                                                                                                                                                                                                              entities requires a license agreement (S or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cercopithecinae;
NCBI_TaxID=36226,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CERMO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CARBOHYD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       | | |||||
134 GNDYEDRYY 142
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUBCELIULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH MOLATITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME
                                                                                                                                                                                                                                                                                  SWISS-PROT entry is copyright. It is produced through a ceen the Swiss Institute of Bioinformatics and the EMBL European Bioinformatics Institute. There are no restrict by non-profit institutions as long as its content is
                                                                                                                                                                                                                                                                                                                                                                                                             SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                   (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Similarity
7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 68
60
51
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67
75
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BAJOR PRION PROTEIN.

REMOVED IN MATURE FORM (BY SIMILARITY).

GPI-ANCHOR (BY SIMILARITY).

BY SIMILARITY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
4 X 8 AA TANDEM REPEATS OF P-H-G-G-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    D582B58E2726C99A CRC64;
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                                                                                                                                                                                                                                       (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                 (TME),
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                                                                                                                                                                                                                                                                Usage
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L outstation -
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PROSITE;

PRION_1; 1. PRION_2; 1.

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Best Local
                              Matches
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DISULFID
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REPEAT
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CHAIN
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135 GNDYEDRYY 143
              1 GXDXEDRYY 9
                                                                                                                                                                                    Brain;
                             Similarity 7; Conserv
                                                            246
                              Conservative
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61
69
77
                                                                                                                                                                                    Glycoprotein;
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                                                                   52
68
76
84
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                                                            26900
                                     90.98;
                                                            MW;
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                                     Score 40;
Pred. No.
                                                                                                                N-LINKED
5 X 8 AA
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GPI-ANCHOR (BY SIMILARITY).
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MAJOR PRION PR
                                                                                                                               N-LINKED
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                                                            835D147CA2B4FDD3 CRC64;
                             Mismatches
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KED (GLCNAC. . .) (POTENTIAL).
AA TANDEM REPEATS OF P-H-G-G-G-W-G-
                                    DB 1;
0.31;
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                              2
                                            Length 246;
                             Indels
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0

Search completed: March Job time : 6.4375 secs 24, 2003, 17:20:17

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Scoring table:
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Maximum DB seq length: 2000000000
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
                                                                                                                                                                                           Score
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94.193 Million cell updates/sec
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Listing first 45 su
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sp_unclassified:*
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097912
097904
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Q8TBG0
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90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9	90.9
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ALIGNMENTS

RESULT 1 097905

097905;

PRELIMINARY;

141 AA

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                              Query Match 93.2%; Score 41; DB 6; Length 141; Best Local Similarity 77.8%; Pred. No. 0.65; Matches 7; Conservative 0; Mismatches 2; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                        Tursiops truncatus (Atlantic bottle-nosed dolphin).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Cetacea; Odontoceti; Delphinidae;
                                                                                                                    NON_TER
                                                                                                                                                                                                                                 MEDIJINE-9303687; PubMed-10373359;
MEDIJINE-9303687; PubMed-10373359;
Mopfnor F., Weidenhofer G., Schneider R., von Brunn A., Gilch S.,
Schwarz T.F., Werner T., Schatzl H.M.;
Rohwarz T.F., Werner T., Schatzl H.M.;
"Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein.";
J. Mol. 18101. 289:1163-1178(1999).
EMBL; AF117311; AAD19992.1;
EMBL; AF117311; AAD19992.1;
HSSP; P10279; 1DWY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prion protein (Fragment).
                                                                                                                                                    Pfam; PF00377; prion; 1.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9739;
                                                                                                                                                                                                                      InterPro: IPR000817; Prion.
   1 GXDXEDRYY 9
                                                                                                      141
141 AA;
                                                                                                       15066 MW;
                                                                                                       79BE306E2AAA187C CRC64;
                                          0;
                                          Gaps
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98

GSDYEDRYY 106

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O97911

DDT C

DDT C

DDT C

OOC C

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RA HOpfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S.,

RA Schwarz T.F., Werner T., Schatzl H.M.;

RT "Analysis of 27 mammalian and 9 avian PrPs reveals high conservation

RT of flexible regions of the prion protein.";

LJ Mol. Biol. 289:1163-1178(1999).

DR EMBL; AF117326; AAD1997.1; ".

DR HSSP; P10279; 1DWY.

DR InterPro; IPR002395; Kininogen.

DR InterPro; IPR002395; Kininogen.

DR InterPro; IPR0017; prion; i.

DR PRINTS; PR0034; KININOGEN.

DR PRINTS; PR00341; PRION.

DR PRINTS; PR00314; PRION.

DR PRINTS; SM00157; PRP; 1.

DR PROSITE; P8001291; PRION_1; 1.
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         MEDIINE-9303687; PubMed-10373359;
Mopfner F., Weldenhofer G., Schneider R., von Brunn A., Gilch S.,
Schwarz T.F., Werner T., Schatzl H.M.,
Analysis of 27 mammalian and 9 avian PrPs reveals high conservation
of flexible regions of the prion protein.";
J. Mol. Biol., 289:1163-1178(1999).
                                                                                                                                                                                                                                                                                                   Bison bonasus (European bison).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovinae; Bison.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    O97912 PRELIMINARY; PRT; 200 AA. 097912; O1-MAY-1999 (TrEMBLrel. 10, Created) O1-MAY-1999 (TrEMBLrel. 10, Last sequence update) O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                             NCBI_TaxID-9902;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prion protein (Fragment).
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01-MAY-1999 (TIEMBLIEL 10, Last sequence update)
01-JUN-2002 (TIEMBLIEL 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea; Bovoidea; Caprinae; Budorcas.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     097911
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Budorcas taxicolor (takin).
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les 7; Conserv
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AF117328; AAD19999.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      181
181 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              93.2%;
77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   19253 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Score 41; DB 6; Length 181; Pred. No. 0.84;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               2;
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                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Interpro; IPR002395; Kininogen.
Interpro; IPR001610; PAC.
Interpro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
PRINTS; PR0034; KININOGEN.
PRINTS; PR00334; PRION.
SMART; SM00086; PAC; 1.
SMART; SM000157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
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InterPro; IPR001817; Prion.
Pfam; PF00377; Prion; 1.
PRINTS; PR00334; KININGGEN.
PRINTS; PR00341; PRION.
SMART; SM00086; PAC; 1.
SMART; SM00186; PAC; 1.
SMART; SM00187; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
NON_TER
200 200
                                                                                                                                                                                                                                                                                                                                         NON_TER
   118
                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDIINE-99303687; PubMed-10373359; Mopfner F., Weidenbefer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Wenner T., Schatzl H., Schwarz T.F., Wenner T., Schatzl H., Schwarz T.F., Wenner T., Schatzl H., Schwarz T.F., Wolley G., W
                                                                                                                                                                                                                                                                                                                                                                                                    PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 J. Mol. Biol. 289:1163-1178(19
EMBL; AF117310; AAD19981.1; -.
HSSP; P10279; 1DWY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       097904;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-MAY-1999 (TrEMBLrel 10,
01-MAY-1999 (TrEMBLrel 10,
01-JUN-2002 (TrEMBLrel 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Prion protein (Fragment).
                                                                                                                                             Local Similarity nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
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InterPro; IPR002395; Kininogen.
                                                                      1 GXDXEDRYY 9
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GSDYEDRYY 126
                                                                                                                                                                                                                                                                                                  215 AA;
                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                     215
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200 AA;
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                                                                                                                                                                                                                                                                                          23182 MW;
                                                                                                                                                             93.2%; Score 41; DB 6; Length 215; 77.8%; Pred. No. 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                93.2%;
77.8%;
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Last annotation update)
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Pred. No. 0.94;
                                                                                                                                                                                                                                                                       97A36721B1E73AE6 CRC64;
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                                                                                                                                     Mismatches
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                                                                                                                         0;
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                                                                                                                     Gaps
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RESULT

P78446

PRELIMINARY;

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RESULT 6
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    RRRR ROCCOGNE DT TO
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Best Local
                                                                                                                                                                                                                                                                             Q15216 PRELIMINARY;
Q15212;
Q1-202 (TrEMBLrel. 01,
01-NOV-1996 (TrEMBLrel. 01,
01-JUN-2002 (TrEMBLrel. 21,
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01-MAY-1997
01-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      VARIANT
NON_TER
                                                                                                                                                                Homo sapiens (Human).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Proc. Natl. Acad. Sci. U.S.A. 88:10926-10930(1991).
EMBL; S71208; AAB20521.1; -
EMBL; M81930; AAB59443.1; -
EMBL; S71212; AAB20523.1; -
EMBL; S71210; AAB20523.1; -
EMBL; S71210; AAB20522.1; -
EMBL; S71210; AAB20522.1; -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE-93250789; PubMed-1363802;
Diedrich J.F., Knopman D.S., List J.F., Olson
Emory C.R., Sung J.H., Haase A.T.;
*Deletion in the prion protein gene in a demen
Hum. Mol. Genet. 1:443-444(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                      MEDLINE-91328137; PubMed-1678248; Puckett C., Concannon P., Casey C.
                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                        PRP OR PRNP.
                                                                                                                                                                                                                                                            Prion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Pfam; PF00377; prion; 1. PRINTS; PR00341; PRION. SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                gene."
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Goldfarb L.G., Brown P., McCombie W.R., Goldgaber D., Swergol Wills P.R., Cervenakova L., Baron H., Gibbs C.J. Jr., Gajdusc "Transmissible familial Creutzfeldt-Jakob disease associated"
                                                                     TISSUE-BRAIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDLINE-92073400; PubMed-1683708;
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                                                                                                                                            NCBI_TaxID-9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NCBI_TaxID=9606;
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    Genomic
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               7; Conserv
2., Concannon P., Casey structure of the human
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233
233 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
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233
25442 MW;
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Last annotation updat
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Last annotation updat
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                                                                                                                                                              Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                  PRT;
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prion
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protein gene.";
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           patient.";
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         "A prion-linked psychiatric disorder.";
Nature 390:241-241(1997).
EMBL; AF030575; AAC05365.1; -.
HSSP; P04156; 1QLZ.
InterPro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Samaia H.B., M
Brentani R.R.;
                                                                                                                              SEQUENCE FROM N.A. MEDLINE-98044028;
                                                                                                                                                                     Homo sapiens (Human).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                    01-AUG-1998 (TrEMBLrel. 07,
01-AUG-1998 (TrEMBLrel. 07,
01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                       060489
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Pfam; PF00377; prion; 1
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                          Prion protein
                                                                                                                                                                                                                                                                           060489;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Perry R.T., Go R.C., Harrell L.E., Acton R.T.;
"SSCP analysis and sequencing of the human prion protein gene (PRNP)
detects two different 24 bp deletions in an atypical Alzheimer's
disease family.";
                                                                                                                                                                  NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE OF 41-85 FROM N.A. MEDLINE=96090306; PubMed=7485229;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Diedrich J.F., Knopman D.S., List J.F., Olson K., Frey W.H., Emory C.R., Sung J.H., Haase A.T.;
"Deletion in the prion protein gene in a demented patient.";
"Lum. Mol. Genet. 1:443-444(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Lee I.Y., Westaway D., Smit A.F., Wang K., Cooper C., Yao H., Prusiner S.B., Hood L., Submitted (FEB-1998) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE OF 9-232 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE=93250789; PubMed=1363802;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                134 GSDYEDRYY 142
                                                                                                                                                                                                                                                                                                                                                                      1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                         ocal
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                                                                                                                                                                                                                                                                                                                                                                                              Similarity 7; Conserv
PS00291; PRION_1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                          245 AA;
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                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                      PRELIMINARY;
                                                                                                                 Mari J.J., Vallada
                                                                                                                                                                                                                        TrEMBLrel. 07, Last sequence update)
TrEMBLrel. 21, Last annotation update)
Variant (Fragment).
                                                                                                                              PubMed=9384372;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            60:12-18(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                          26884 MW;
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                                                                                                                                                                                                                                                              Created)
                                                                                                                                                                                                                                                                                                                                                                                           Score 41; DB 4;
Pred. No. 1.2;
0; Mismatches
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                                                                                                                 H.P.,
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                                                                                                                 R.P.,
                                                                                                                                                                                                                                                                                                                                                                                                                  Length 245;
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                                                                                                                 А.J.,
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RESULT
OUT-19
ID UT-19
ID UT-1
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Q96E70
           RRR ROCC OCCUPATO
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Best Local Similarity
Thes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
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NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                  Q96E70 PRELIMINARY; PRT; 253 AA. Q96E70; Q96E70; Q1-DEC-2001 (TrEMBLrel. 19, Created) 01-DEC-2001 (TrEMBLrel. 19, Last sequence update) 01-MAR-2002 (TrEMBLrel. 20, Last annotation update) 51milar to prion protein (p27-30) (Creutzfeld-Jakob Gerstmann-Strausler-Scheinker syndrome, fatal famili Homo saplens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-20049886; PubMed-10581485;
Cervenakova L., Buetefisch C., Lee H.S., Taller I., Stone Gibbs C.J. Jr., Brown P., Hallett M., Goldfarb L.G.;
"Novel PRNP sequence variant associated with familial encephalopathy.";
Am. J. Med. Genet. 88:653-656(1999).
EMBL; AF076976; AAD46098.1; -.
HSSP; P04156; 1QLZ.
InterPro; IPR00017; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q9UP19;
01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-JUN-2002 (TrEMBLrel. 21,
   Strausberg R.;
Submitted (AUG-2001)
                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                            Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00377; prion; 1
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
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Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                            TISSUE-OVARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
VARIANT 187 187
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Q9UP19
                                                                                                                                                                                          NCBI_TaxID=9606;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  142 GSDYEDRYY 150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Similarity 77.8
7; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PS00706; PRION_2; 1.
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                                                                                                                                                                                                                        Chordata; Craniata; Vertebrata; Euteleostomi; Primates; Catarrhini; Hominidae; Homo.
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77.88;
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Last annotation update)
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       EMBL/GenBank/DDBJ databases
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Pred. No. 1.2;
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1.2;
                                                                                                                                                                                                                                                                                                                                          fatal familial insomnia).
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                                                                                                                                                                                                                                                                                                                                                                             disease,
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Query Match
Best Local Similarity
Matches 7; Conser
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Best Local
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                                                                                                                                                                                                                                                                                                                                                         Q95M08 PRELIMINARY;
Q95M08;
01-DEC-2001 (TrEMBLrel. 1
01-DEC-2001 (TrEMBLrel. 1
01-MAR-2002 (TrEMBLrel. 2
Matsumoto Y., Endo H., Onodera T.;

Matsumoto Y., Endo H., Onodera T.;

"Comparative analysis of the prion protein ORF nucleotide from two wild ruminants, moufflon and golden takin.";

Submitted (APR-2001) to the EMBL/GenBank/DDBJ databases.

EMBL; AB060290; BAB69957.1; -

InterPro; IPR000817; Prion.

Pfam; PF00377; prion; 1.

PROSITE; PS00291; PRION_1; UNKNOWN_1.
                                                                                                                                                         Seo S., Hara K., Kubosaki A., Nasu Y., Nishimura Matsumoto Y., Endo H., Onodera T.;
                                                                                                                                                                                                                                               Budorcas taxicolor (takin).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eute
Mammalla; Eutheria; Cetartiodactyla; Ruminantia; Pecora;
Bovidae; Caprinae; Budorcas.
                                                                                                                                                                                                                                                                                                                          Prion protein. PRNP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Strausberg R.;
Submitted (FEB-2002) to the EMBL/GenBank/DDBJ databases
EMBL; BC022532; AAH22532.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-JUN-2002 (TrEMBLrel. 21, Created)
01-JUN-2002 (TrEMBLrel. 21, Last sequence up
01-JUN-2002 (TrEMBLrel. 21, Last annotation
Hypothetical 27.6 kDa protein.
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InterPro; IPR000817; Prion.
Pfam; PF00377; Prion; 1.
PROSITE; PS00291; PRION_1; UNKNOWN_1.
PROSITE; PS00706; PRION_2; UNKNOWN_1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Hypothetical SEQUENCE 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Primates;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Q8TBG0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         OSTBG0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE
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                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                              NCBI_TaxID=37181;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TISSUE-BRAIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (Human)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NCBI_TaxID=9606;
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142 GSDYEDRYY 150
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity 77.177; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            253 AA; 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   253 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .27629 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     27629 MW; 43DB4D3BA5F67F84 CRC64;
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77.8%;
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77.8%;

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0
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Pred. No. 1.2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              43DB569A1FF67F84 CRC64;
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1.2;
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                                                                                                                                                                        Τ.,
                                                                                                                                                                                                                                                                   Euteleostomi;
cora; Bovoidea;
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RESULT 13
Q9TU20
ID Q9TU2
AC Q9TU2
DT 01-MJ
DT 01-MJ
DT P110-
GN PRP.
OS VBTC
CC MANM
OC MANM
RN (11)
RP SEQ
RP SEQ
RP SEQ
RT *UT
RT SUE
SUE
DR HS
DR HS
DR PF
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Interpro; IPRO00817; prion.
Priam; pr00377; prion; 1.
Print; pr00377; prion; 1.
PRINTS; pr00341; prioN.
SMART; SM00157; prR; 1.
PROSITE; pr00705; prION_1; 1.
PROSITE; pr00706; prION_2; 1.
SEQUENCE 285 AA; 30769 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            075942
075942;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cervenskovs L., Halperin J.J., Stone G., Taller I., Gibbs C.J. Jr.; *A new Creutzfeldt-Jakob disease family with 96bp insertion mutation in the PRNP gene."; Submitted (NOV-2000) to the EMBL/GenBank/DDBJ databases. EMBL; AF085477; AAC62750.2; -. HSSP; P04156; 10LZ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-NOV-1998 (TrEMBLrel. 08, 01-MAR-2001 (TrEMBLrel. 16, 01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID-9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              145 GSDYEDRYY 153
  Gilch S., Schatzl H.M.;
"Unusual prion protein octarepeat structure of the highly BSE-
susceptible lemur monkey.";
Submitted (AUG-1999) to the EMBL/GenBank/DDBJ databases.
EMBL; AF177293; AAD54335.1; -.
HSSP; P04925; 1AG2.
InterPro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
PRINTS; PR00341; PRION.
                                                                                                                                                                                                                                                                                                                             Varecia variegata variegata.
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Strepsirhini; Lemuridae; Varecia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAY-2000 (TrEMBLrel. 13,
01-MAY-2000 (TrEMBLrel. 13,
01-DEC-2001 (TrEMBLrel. 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Q9TU20;
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                                                                                                                                                                                                                                                                                                                                                                                                                                      Prion protein (Fragment).
                                                                                                                                                                                                                        SEQUENCE FROM N.A.
G11ch S., Schatzl H.M.;
                                                                                                                                                                                                                                                                                                   NCBI_TaxID=87289
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     174 GSDYEDRYY 182
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Local Similarity
les 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      1 GXDXEDRYY 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27860 MW;
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77.88;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       93.2%;
77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Last sequence update)
Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Created)
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Last annotation update)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 C14244BB1183F653 CRC64;
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O97903 PRELIMINARY; PRT; 195 AA.

O97903;
O1-MAY-1999 (TrEMBLrel. 10, Created)
O1-MAY-1999 (TrEMBLrel. 10, Last sequence update)
O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)

195 AA

RESULT 15

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RESULT 14
097694
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                                                                                                                                 Query Match
Best Local :
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InterPro; IPR000817; Prion.
Pfam; PF00377; Prion; 1.
PRINTS; PR0034; KININOGEN.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Cervoldea; Cervidae; Cervinae; Cervus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          01-MAY-1999 (TREMBLICEL 10, 01-MAY-1999 (TREMBLICEL 10, 01-JUN-2002 (TREMBLICEL 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schatzl H.M.; Schwarz T.F., Werner T., Schatzl H.M.; Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein."; Mol. Biol. 289:1163-1178(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prion protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             097694;
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                                                                                                                                                                                                                                        NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; AF113941; AAD13289.1; -. HSSP; P10279; 1DWY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE FROM N.A. MEDLINE-99303687; PubMed=10373359;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       animals?":
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Schatzl H.M., Wopfner F., Gilch S., von Brunn "Is codon 129 of prion protein polymorphic in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A. MEDLINE-97317556; PubMed-9174569;
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                                                                                                                                                                                                               SEQUENCE
                                                                                                                                                                                                                                                                                                  PROSITE; PS00291; PRION_1; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         75 GNDYEDRYY 83
121 GNDYEDRYY 129
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                                                                                                    Local Similarity
les 7; Conserv
                                                   1 GXDXEDRYY 9
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                                                                                                                                                                                                                 185 AA; 19870 MW; BB87C7658BC66E79 CRC64;
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                                                                                                       Conservative
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: 13436 MW; CC2C8A5A855A7C94 CRC64;
                                                                                                                                                                                                                                              185
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                                                                                                          90.9%; Score 40; DB 6; Length 185; 77.8%; Pred. No. 1.4; tive 0; Mismatches 2; Indels
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Pred. No. 0.91;
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human beings but not in
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Prion protein (Fragment).

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Search completed: March 24, 2003, 17:22:11 Job time: 20.6875 secs
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                                                                                                   Query Match
Best Local Similarity
""" conserve
                                                                                                                                                                                                    RX MEDLINE-9303687; PubMed-10373359;
RA WEDLINE-9303687; PubMed-10373359;
RA WOpfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S.,
RA Schwarz T.F., Weiner T., Schatzl H.M.;
RT Analysis of 7 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein.";
RT J. Mol. Biol. 289:1163-1178(1999).
RE EMBL; AFI17309; AAD19980.1; -.
DR HSSP; P10279; IDMY.
DR HSSP; P10279; IDMY.
DR InterPro: IPR002395; Kininogen.
DR FAINTS; PR000817; Prion; 1.
DR Pfam; PF00377; prion; 1.
DR PRINTS; PR00341; PRION.
DR PRINTS; PR00341; PRION.
DR SMART; SM00157; PRP; 1.
DR PROSITE; PS00291; PRION_1; 1.
DR PROSITE; PS00291; PRION_2; 1.
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SEQUENCE
                                                           109 GNDYEDRYY 117
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Hippotraginae; Addax.
NCBI_TaxID=59515;
[1]
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE FROM N.A.
                                                                            1 GXDXEDRYY 9
                                                                                                                                                                           195
195 AA;
                                                                                                                 90.9%; Score 40; DB 6; Length 195; Length 195; Conservative 0; Mismatches 2; Indels
                                                                                                                                                   195; 21321 MW; 6A9BA6A7ElAFECA9 CRC64;
                                                                                                                0; Mismatches
                                                                                                           Indels
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                                                                                                         Gaps
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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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seq length: 2000000000
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Match
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                         Issued_Patents_AA: *
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/cgn2_6/ptodata/2/laa/5B_COMB.pep:*
/cgn2_6/ptodata/2/laa/6A_COMB.pep:*
/cgn2_6/ptodata/2/laa/6B_COMB.pep:*
/cgn2_6/ptodata/2/laa/BCTUS_COMB.pep:*
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US-08-244-701B-43
US-08-244-701B-10
US-08-244-701B-12
US-09-076-721-10
US-09-076-721-10
US-09-076-721-11
US-09-076-721-2
US-09-076-721-7
US-09-076-721-7
US-09-076-721-7
US-09-076-721-9
US-08-556-823-10
US-08-556-823-10
US-08-560-626-8
US-08-59-261A-2
US-08-59-261A-2
US-08-59-261A-2
US-08-59-261A-2
US-08-59-261A-2
US-08-59-261A-2
US-08-59-261A-2
US-08-59-261A-3
US-09-128-450-20
US-09-128-450-20
US-09-823-494-20
US-09-550-374-2
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US-08-560-626-9
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Result

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2254 2254 2254 2254 2254 2254

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Database :

Minimum DB Maximum DB

Scoring table: Sequence:

Perfect score:

OM protein -

ALIGNMENTS

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US-08-244-701B-43
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TELEFAX: (214)
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 43
SEQUENCE CHARACTERISTICS:
FENGTH: 21 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 43, Application US/08244701B Patent No. 5773572
GENERAL INFORMATION:
                        1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0.
                                                                                                                                                                                                                                                                                                    NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 800
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STREET: LILL KITTE: New York York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: Fishleigh, Rol
APPLICANT: Robson, Barry
APPLICANT: Mee, Roger P.
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CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TITLE OF INVENTION: Fraq
                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STATE: New York COUNTRY: U.S.A.
                                                               Local Similarity
nes 7; Conserv
                                                                                                                                                                                TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/0
FILING DATE: 02-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
 11 GSDYEDRYY 19
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                     (212) 869-8864/9741
                                                               Conservative
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                                                                                                                                                  peptide
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                                                                               93.2%;
77.8%;
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                                                                 0;
                                                               Score 41; DB 1;
Pred. No. 0.087;
0; Mismatches
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                                                                                               Length 21;
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US-09-076-721-43
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                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE CHARACTERISTICS:
LENGTH: 21 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                          NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REGISTRATION NUMBER: 30,256
REFERENCE, DOCKET NUMBER: 807
TELECOMMUNICATION INFORMATION:
                                                                                                        TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/076,721
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COMPUTER READABLE FORM:
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APPLICATION NUMBER:
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CORRESPONDENCE ADDRESS:
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   CITY: New York STATE: New York COUNTRY: U.S.A.
                             ADDRESSEE: Pennie o Eumanie...
STREET: 1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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ZIP: 10036
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Fanucci, Allan A.
Fanucci, Allan A.
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Mee, Roger P.
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                                                                                                                                    Barry
                                                                                            Fragments of Prion Proteins 67
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Pred. No. 0.087;
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US-08-244-701B-12
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                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                      APPLICANT:
APPLICANT:
                     COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                    COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                          NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                         TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MOLECULE TYPE:
            APPLICATION NUMBER: US/08/244,701B
                                                                                                                                COUNTRY:
                                                                                                                                                    STATE:
                                                                                                                                                              CITY: New York
                                                                                                                                                                              STREET:
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LOCATION:
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                                                                                                                                                  New York
                                                                                                                                                                              1155 Avenue of the Americas
                                                                                                                                                                                                                                                  Mee, Roger P.
                                                                                                                                   U.S.A.
                                                                                                                                                                                                                                                                                 Fishleigh,
                                                                                                                                                                                                                                                                 Robson, Barry
                                                                                                                                                                                             Pennie & Edmonds
                                                                                    Floppy disk
02-JUN-1994
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Fragments of Prion Proteins

Robert V.

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INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
FILING DATE: 02-JUN-199
CLASSIFICATION: 436
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REFERENCE/DOCKET NUMBER: 80 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,
                                                                                                                                          OTHER INFORMATION:
OTHER INFORMATION:
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MOT.EX: 66141 PENNIE
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OPERATING SYSTEM:
12 GSDYEDRYY 20
                                                                                                                                                                                                                                                        OTHER INFORMATION:
                                                            Local Similarity
les 7; Conserv
                                                                                                                                                                                        NAME/KEY: Modified-site
                                                                                                                                                                                                                                           OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (212) 790-9090
(212) 869-8864/9741
                                                            Conservative
                                                                                                                                                                                                                                                                                                        Modified-site
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PatentIn Release #1.0, Version #1.25
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               IBM PC compatible
SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                         linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Floppy disk
                                                                                                                                                                                                                                                                                                                                        peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  02-JUN-1994
                                                                            93.2%;
                                                                                                                                                                                                                                    /label= x
/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
                                                                                                                                         /label= Y
/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US/08/244,701B
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                                                                                       Score 41;
                                                                          Pred. No.
                                                                                         DB 1;
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                                                                                         Length 25;
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Patent No.
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Best Local !
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SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT:
APPLICANT:
                                                                                                                                                                                  COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                    NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
                                                                                                APPLICATION NUMBER:
                                                                 ATTORNEY/AGENT INFORMATION:
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NAME: Fanuci, Allan A.
REGISTRATION UNMBER: 30,256
REFERENCE/DOCKET NUMBER: 8080-007
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: Modified-site LCCATION: 25
OTHER INFORMATION: /label
OTHER INFORMATION: /note-
OTHER INFORMATION: of X a
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ELECOMMUNICATION INFORMATION:
            NAME: Fanucci, Allan A. REGISTRATION NUMBER: 30,256 REFERENCE/DOCKET NUMBER: 808
                                                                                                                                                                                                                                                                                                                                                         STREET:
                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                       COUNTRY:
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                                                                                                                                                                   APPLICATION NUMBER:
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OTHER INFORMATION:
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6379905
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Mee, Roger P.
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(212) 869-8864/9741
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linear
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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                8080-007
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Pred. No. 0
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RESULT 6
US-09-076-721-12
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                                                          APPLICATE:
FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: FANUCCI, Allan A.
REGISTRATION NUMBER: 30,256
REGISTRATION NUMBER: 8080.
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION 1690-9090
TELECOMMUNICATION 1690-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Patent No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 12,
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                                INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (212) 869-886
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
SEQUENCE CHARACTERISTICS:
LENGTH: 25 amino acids
                                                                                                                                                                                                       CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Robson, Barry
APPLICANT: Mee, Roger P.
TITLE OF INVENTION: Fragments of Prion Proteins
NUMBER OF SEQUENCES: 67
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                                                                                                                                                                                                                                                                                                                                                                                                   STATE: New York COUNTRY: U.S.A. ZIP: 10036
                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: New York
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LOCATION:
                                                                                                                                                                                                                                                            FILING DATE:
                                                                                                                                                                                                                                                                          APPLICATION NUMBER:
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ues 7; Conserv
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OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         LENGTH: 25 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELEPHONE:
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6379905
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1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   : (212) 790-9090
(212) 869-8864/9741
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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Pred. No. 0.1;
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                                   Query Match
     Matches
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                                                                                                                                                                                TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                         MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PEATURE: Modified-site
                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 26 amino acids
                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 800
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TITLE OF INVENTION: Fragments of Prion Proteins NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
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OTHER INFORMATION: /label
OTHER INFORMATION: /notee
OTHER INFORMATION: of X 8
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MOLECULE TYPE: peptide
                                                                               MOLECULE TYPE:
 Local Similarity 77.1 hes 7; Conservative
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Local Similarity 77.88;
es 7; Conservation
                                                                                                               STRANDEDNESS:
                                                                                                                              TYPE: amino acid
                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/0 FILING DATE: 02-JUN-1994 CLASSIFICATION: 436
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OTHER INFORMATION:
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STRANDEDNESS: single
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                                                                                peptide
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               93.2%;
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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of Y and denotes one or more amino acid(s)"
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Score 41; DB 1 pred. No. 0.11; 0; Mismatches
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                                 DB 1; Length 26;
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     Indels
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US-09-076-721-41
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                                                                                                  Sequence 7, Application US/08244701B Patent No. 5773572 GENERAL INFORMATION:
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Best Local :
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICANT: Robson, F
APPLICANT: Mee, Roge
TITLE OF INVENTION:
                                                         APPLICANT:
APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
    NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TELECOMMUNICATION INFORMATION: TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CORRESPONDENCE ADDRESS:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                TYPE: amino ac
STRANDEDNESS:
TOPOLOGY: line
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STREET: 1155 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          10 GSDYEDRYY 18
                                                                                                                                                                                                       10 GSDYEDRYY 18
                                                                                                                                                                                                                                                             Local Similarity
nes 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Fanucci, Allan A. REGISTRATION NUMBER: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/076,721 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CLASSIFICATION:
 ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICATION NUMBER:
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10036
                                                                                                                                                                                                                                                                                                                                                                           H: 26 amino acids amino acid
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                                                                      Robson,
                                                                                                                                                                                                                                                              Conservative
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                                                                                    Fishleigh, Robert V.
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linear
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Pennie & Edmonds
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                                                                     Barry
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                                                                                                                                                                                                                                                                            93.2%;
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Pred. No. 0.11;
0; Mismatches
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COUNTRY:

U.S.A.

New York New York

1155 Avenue of the Americas

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RESULT 10
US-08-244-701B-9
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LORPY disk

OPERATING SYSTEM: PC-DOS/MS-DOS

SOFTWARE: Patentin Release #1.0, Version #1.25

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/244 7017

FILING DATE: 02-TIN-12
                                                                                                                                                                                                                                                                           Sequence 9, Application US/08244701B Patent No. 5773572
                                                                                                                                                                                                                                                                                                                                                                                                                                               Matches
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NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 8080
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
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TELEFAX: (212) 869-8864/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 7:
                                                                                                                                                                                                                                                               GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE CHARACTERISTICS:
LENGTH: 31 amino acids
TYPE: amino acid
STRANDEDNESS: single
                        COUNTRY: U.S.A.
ZIP: 10036
ZIP: 10036
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                             NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                        TITLE OF INVENTION:
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                                                                                                                                                                                                               APPLICANT:
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OTHER INFORMATION:
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hes 7; Conserv
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OTHER INFORMATION:
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                                                                                              STREET: 1155 A
CITY: New York
STATE: New Yor
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COMPUTER: IBM PC compatible OPERATING SYSTEM: PC-DOS/MS-DOS
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                                                                                              New York
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linear
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                                                                                                                                                                                Roger P.
ON: Fragments of Prion Proteins
CES: 67
                                                                                                                                                                                                                                Barry
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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/note= "X may be absent or present independently of Y and denotes one or more amino acid(s)"
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Patent No. 6379905
GENERAL INFORMATION:
APPLICANT: Fishleigh, Robert V.
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OLECUI.

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ACTERISTIC
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APPLICATION NUMBER: US,
FILING DATE: 02-JUN-199
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                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CORRESPONDENCE ADDRESS:
                                                                  PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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ATTORNEY/AGENT INFORMATION:
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OTHER INFORMATION: /note= "Y may be absent or present independently
OTHER INFORMATION: of X and denotes one or more amino acid(s)"
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LOCATION:
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les 7; Conserv
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                                                                                                                                        CLASSIFICATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET:
                                    FILING DATE
                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                       APPLICATION NUMBER: US/09/076,721
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                  10036
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      : (212) 790-9090
(212) 869-8864/9741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mee, Roger P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Robson, Barry
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77.88;
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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                                                                      US 08/244,701
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Pred. No. 0.13;
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Gaps

TELECOMMUNICATION INFORMATION:

REGISTRATION NUMBER: 30 REFERENCE/DOCKET NUMBER:

8080-007

Fanucci, Allan A.

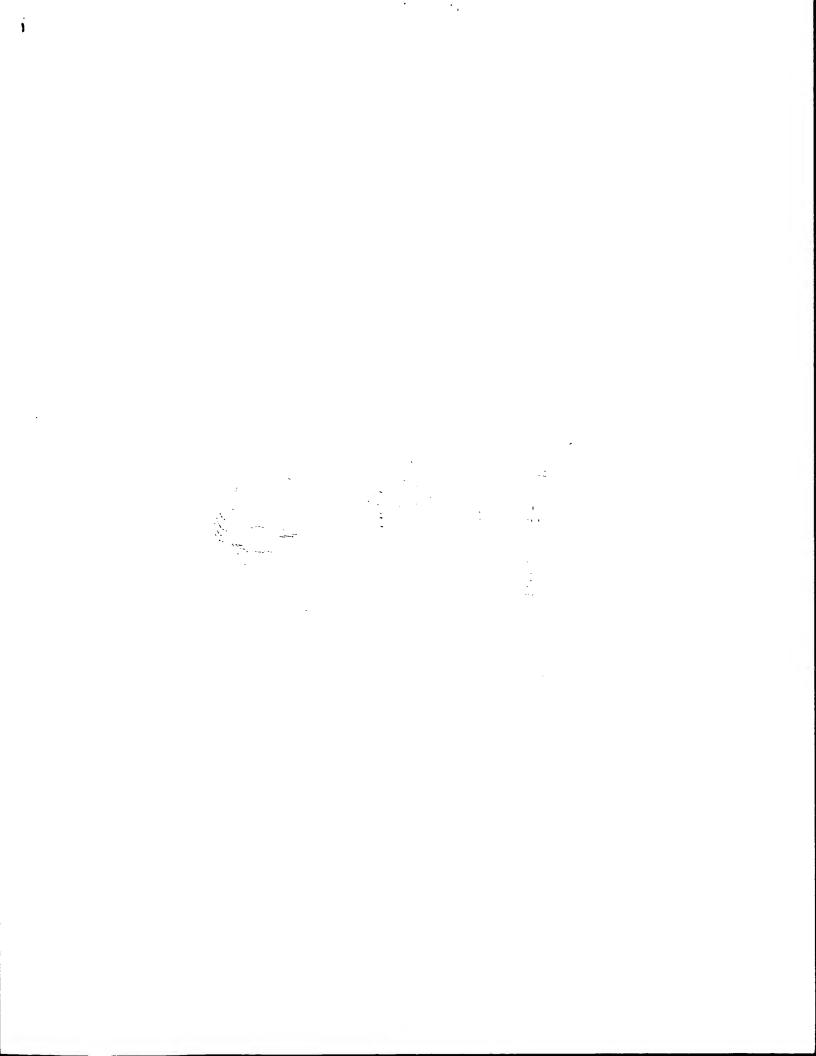
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                            ATTORNEY/AGENT INFORMATION:
NAME: FADUCCI, Allan A.
REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 806
TELECOMMUNICATION INFORMATION:
                                                                                                                             APPLICATION NUMBER:
                                                                                                                                                                                                CURRENT APPLICATION DATA: APPLICATION NUMBER: US,
                                                                                                                                                                                                                                                                           MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                          CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Robson, Barry
APPLICANT: Mee, Roger P.
TITLE OF INVENTION: Fragn
                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
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                                                                                                                  FILING DATE:
                                                                                                                                                                                                                        COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
                  TELEPHONE:
                                                                                                                                                                CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                            STREET: 1155 Avenue of the Americas CITY: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                              COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                             ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       12 GSDYEDRYY 20
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tes 7; Conserv
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OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                10036
                                                                                                                                                                                                                                                                                                                                             New York
: (212) 790-9090
(212) 869-8864/9741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Fishleigh, Robert V.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
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(212) 869-8864/9741
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
                                                                                                                               US 08/244,701
                                                                                                                                                                                                US/09/076,721
                                               8080-007
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Pred. No. 0.
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; TOPOLOGY: 1:
; MOLECULE TYPE:
US-08-556-823-10
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US-08-556-823-10
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Patent No.
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                                                                                            TELEPHONE: (415) 322-
TELEFAX: (415) 854-08
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 31 amino acids
                                                          SEQUENCE CHARACTERISTICS:
LENGTH: 142 amino acids
                                                                                                                                       ATTORNEY/AGENT INFORMATION:
NAME: VALETA Greg
REGISTRATION NUMBER: 35,127
REFERENCE/DOCKET NUMBER: 07532/003001
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                       SOFTWARE: ASCIII CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                       COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy
COMPUTER: IBM PC com
                                                                                                                                                                                                                                                                                                                                                                                                                                                            NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
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                                                  TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LENGTH: 31 amino a
TYPE: amino acid
STRANDEDNESS: since
                                                                                                                                                                                                                    FILING DATE:
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                   MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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OTHER INFORMATION: /note= "Y may be absent or present independently
OTHER INFORMATION: of X and denotes one or more amino acid(s)"
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                                                                                                                                                                                                                                                    APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                        STATE:
                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
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5750361
                                             amino acid
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                                                                                                           (415) 322-50
(415) 854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Stanley B. Prusiner
Kiyotoshi Kaneko
Fred E. Cohen
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                               linear
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               peptide
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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Pred. No. 0.13;
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US-08-242-188-2
                                                                                                                                                               US-08-509-261A-2
                                                                                                                                                                                                                                                                                                                                                                                                                 REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06510/014001
TELECOMMUNICATION: INFORMATION:
TELEPHONE: (415) 854-5277
TELEPHONE: (415) 854-6875
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 253 amino acids
TYPE: beningle
TOPOLOGY: linear
MOLECULE TYPE: peptide
ORIGINAL SOURCE:
ORGANISM: HUMAN PRION PROTEIN, HuPrP
                                                                                                  Sequence 2, Application US/08509261A Patent No. 5763244 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence 2, Application US/08242188 Patent No. 5565186
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GENERAL INFORMATION:
APPLICANT: Prusiner, Stanley B.
APPLICANT: Scott, Michael R.
APPLICANT: Telling, Glenn
APPLICANT: Telling, Glenn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/242,188
FILING DATE: 13-MAY-1994
CLASSIFICATION: 435
ATTORNEY/AGENT INFORMATION:
NAME: BOSICEVIC, Karl
    APPLICANT: Prusiner, Stanley B. APPLICANT: Scott, Michael R. APPLICANT: Telling, Glenn TITLE OF INVENTION: Method of D TITLE OF INVENTION: in a Sample
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
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ADDRESSEE: Karl Bosicevic
STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION:
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                                                                                                                                                                                                                                             142 GSDYEDRYY 150
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nes 7; Conserv
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77.8%;
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                                                                                                                                                                                                                                                                                                                                         93.2%;
Method of Detecting Prions in a Sample and Transgenic Animal Used fore
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Pred. No. 0.62;
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                                                                                                                                                                                                                                                                                                                                                          Length 253;
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Search completed: March 24, 2003, 17:23:04
Job time: 8.625 secs
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                                                                                                                                                                                                                                                                          INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 253 amino acids
                                                                                                                                                                                                                                                                                                                                                                  ATTORNEY/AGENT INFORMATION:
ATTORNEY/AGENT INFORMATION:
NAME: Bozicevic, Karl
REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 651(
TELECOMMUNICATION INFORMATION:
TELEPHONE: 650-327-3400
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER: US/08/509,261A FILING DATE: 31-UUL-1995 CLASSIFICATION: 800 PRIOR APPLICATION DATA: APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ for Windows Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES: 4
CORRESPONDENCE ADDRESS:
ADDRESSEE: Bozicevic & Reed, LLP
STREET: 285 Hamilton Avenue, Suite 200
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                   142 GSDYEDRYY 150
                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET: 285 Hami
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                                                                                                        1 GXDXEDRYY 9
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                                                                                                                                       Conservative
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                                                                                                                                                         93.2%;
77.8%;
                                                                                                                                                                                                                                                                                                                                                                                                      6510-030001
                                                                                                                                                         Score 41; DB Pred. No. 1.1;
                                                                                                                                       Mismatches
                                                                                                                                                                       DB 1;
                                                                                                                                                                       Length 253;
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Database :
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Searched:
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44
1 GXDXEDRYY 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                             Published_Applications_AA: *
: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US08_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
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: /cgn2_6/ptodata/1/pubpaa/US08_PUBCOMB.pep:*
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: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
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: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US10_PUBCOMB.pep:*
: /cgn2_6/ptodata/1/pubpaa/US60_PUBCOMB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

19	18	17	16	15	14	13	12	11	10	9	8	7	6	5	4	w	2	_	Result
39	39	39	39	39	40	40	40	40	40	40	41	41	41	41	41	41	41	41	Score
88.6	88.6	88.6	88.6	88.6	90.9	90.9	90.9	90.9	90.9	90.9	93.2	93.2	93.2	93.2	93.2	93.2	93.2	93.2	Query Match Length
254	254	254	164	163	256	256	256	256	255	161	264	263	256	253	253	253	162	161	
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US-10-106-574-7	US-10-106-574-6	US-10-106-574-5	US-09-745-003-12	US-09-745-003-11	US-10-109-551-10	US-10-109-551-8	US-10-109-551-6	US-10-109-551-4	US-09-943-906-4	US-09-745-003-7	US-10-209-194-2	US-09-943-906-3	US-10-109-551-2	US-09-943-906-2	US-09-919-172-57	US-09-904-987-3	US-09-745-003-10	US-09-745-003-9	ID
Sequence 7, Appli	Sequence 6, Appli	Sequence 5, Appli	Sequence 12, Appl	Sequence 11, Appl	Sequence 10, Appl	Sequence 8, Appli	Sequence 6, Appli	Sequence 4, Appli	Sequence 4, Appli	Sequence 7, Appli	Sequence 2, Appli	Sequence 3, Appli	Sequence 2, Appli	Sequence 2, Appli	Sequence 57, Appl	Sequence 3, Appli	Sequence 10, Appl	Sequence 9, Appli	Description

TYPE: PRT

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	<u> </u>			· · · ·
RESULT 2 US-09-745-003 ; Sequence 10 ; Patent NO. ; GENERAL INF. ; TITLE OF I ; FILE REFER; CURRENT FI ; CURRENT FI ; NUMBER OF SOFTWARE: ; SEQ ID NO 1 ; FILE REFER; CURGTH: 1	Query M Best Lo Matches Qy 1	RESULT 1 US-09-745-003-9 ; Sequence 9, Applicat patent No. US2002004 ; GENERAL INFORMATION; ; APPLICANT: Bazan, F FILE REFERENCE: PF ; CURRENT APPLICATION ; CURRENT FILING DATE ; NUMBER OF SEQ ID NO ; SEQ ID NO		
THE PART OF TO	Match Local Simila les 7; Co 1 GXDXEDRYY 50 GSDYEDRYY	O9-745-003-9 O9-745-003-9 O9-745-003-9 atent no. US2002004: ENERAL INFORMATION: APPLICANT: Bazan, Frifiz of INVENTION: FILE REFERENCE: PPP: CURRENT APPLICATION CURRENT APPLICATION UNDBER OF SEQ ID NO. SOFTWARE: Patentin' EQ ID NO 9 EQ ID NO 9 EQ ID NO 9 EQ ID NO 9 CORGANISM: bovine O9-745-003-9		22222222222222222222222222222222222222
003-10 10, Application U 10, Application U 10, US20020042122A1 INFORMATION: NT: Bazan, Fernand FF INVENTION: Human FERENCE: prp2 APPLICATION NUMBE FILING DATE: 200 OF SEQ ID NOS: 13 E: Patentin Ver. 2 10 10	similarity 7; Conser %EDRYY 9 	15-003-9 15-003-9 10ce 9, Application US 10ce 9, Application US 10ce 9, Application US 10ce 10ce 10ce 10ce 10ce 10ce 10ce 10ce		55555555555555555555555555555555555555
lion 2122# ernan Huma Huma NUME 20 5: 13	93.2: larity 77.8: Conservative XY 9 	on US/ 122A1 rnando Human NUMBER 2000 : 13 er. 2.		254 439 46 46 46 46 434 46 434 434 803 979 979 979 1938 833 1938 315 315 315 315 315 315 315 315 315 315
709 Pro Pro	8 2 8 8 4;	0 -1 Pr 09		001 001 001 001 001 001 001 001 001
US/09745003 .1 .do J .n Proteins; Related Reagents .neer: US/09/745,003 .00-12-20	Score 41; DB 10; Length 1 Pred. No. 0.35; 0; Mismatches 2; Indels	US/09745003 2A1 ando J man Proteins; Related Reagents MBER: US/09/745,003 2000-12-20 13 . 2.0	ALIGNMENTS	US-10-106-574-8 US-09-943-906-1 US-010-115-984-2 US-09-93-780-4 US-09-966-422B-9 US-09-966-422B-9 US-09-974-298-133 US-09-974-298-133 US-09-9884-457A-4 US-09-9884-001-4 US-09-912-717-3 US-09-912-717-3 US-09-912-717-3 US-09-912-717-3 US-09-912-717-3 US-09-912-717-3 US-09-912-717-3 US-09-912-717-3 US-09-912-717-3 US-10-086-464-2 US-10-086-464-2 US-10-086-464-3 US-09-915-242-12339 US-10-086-464-5 US-09-915-242-12339 US-10-086-464-5 US-09-915-242-12339 US-10-086-464-5 US-09-953-956-2 US-09-953-956-2 US-09-953-956-2
	161; Ls 0;			Sequence 1, Sequence 2, Sequence 4, Sequence 193 Sequence 193 Sequence 4, Sequence 4, Sequence 3, Sequence 3, Sequence 12, Sequence 14, Sequence 14, Sequence 14, Sequence 14, Sequence 15, Sequence 16, Sequence 17, Sequence 17, Sequence 17, Sequence 18, Sequence 19, Sequence 20, Sequence 20, Sequence 21,
	Gaps 0;			1, Appli 4, Appli 4, Appli 4, Appli 9, Appli 193, Appli 494, Appli 494, Appli 494, Appli 2, Appli 1218, Appli 1218, Appli 1218, Appli 1218, Appli 123, Appli 124, Appli 125, Appli 127, Appli 128, Appli 129, App

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                                                                                                                           ; TYPE: PRT
; ORGANISM: Homo sapiens
; FEATURE:
; NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No. US20020119463A1 1256895CD1
US-09-919-172-57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-09-919-172-57
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GENERAL INFORMATION:
APPLICANT: No. US20020037908Alactyl, Inc.
APPLICANT: No. US20020037908Alactyl, Inc.
TITLE OF INVENTION: Methods and Compositions for Controlling Pathological and Prepath
TITLE OF INVENTION: Protain Assembly or Aggregation
FILE REFERENCE: 42108/26146
                                                   Matches
                                                                         Query Match
Best Local :
                                                                                                                                                                                                                                                          NUMBER OF SEQ ID NOS: 102
SOFTWARE: PERL Program
SEQ ID NO 57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 57, Application US/09919172 Patent No. US20020119463A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: PatentIn version 3.0 SEQ ID NO 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 3, Application US/09904987 Patent No. US20020037908A1
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                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/919,172
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: 60/222,469
PRIOR FILING DATE: 2000-07-28
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TYPE: PRT
ORGANISM: homo sapiens
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: NCBI ENTREZ / XM_009567
DATABASE ENTRY DATE: 2001-04-17
RELEVANT RESIDUES: (1)..(253)
                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Turner, Christopher M.
TITLE OF INVENTION: PROSTATE CANCER MARKERS
FILE REFERENCE: PA-0036 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/904,987
CURRENT FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Faris, Mary
APPLICANT: Turner, Chi
                                                                                                                                                                                                                                            LENGTH:
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                                                 Local Similarity les 7; Conserv
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les 7; Conserv
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                                                               93.2%;
77.8%;
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                                           Score 41; DB Pred. No. 0.56 0; Mismatches
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                                                                                 DB 10; Length 253;
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                                           Gaps
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                                           0;
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Best Local Similarity
Thehes 7; Conserv
                                                                                                                                                                                                              RESULT 6
US-10-109-551-2
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US-09-943-906-2
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                                                       Sequence 2, Application US/10109551
publication No. US20020194635A1
GENERAL INFORMATION:
APPLICANT: DUNNE, PATRICK W.
APPLICANT: PIEDRAHITA, JORGE
TITLE OF INVENTION: TRANSGENIC ANIMALS RESISTANT TO TRANSMISSIBLE
TITLE OF INVENTION: SPONGIFORM ENCEPHALOPATHIES
FILE REFERENCE: TAMK: 207US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 2, Application US/09943906 Patent No. US20020150571A1 GENERAL INFORMATION:
CURRENT APPLICATION NUMBER: US/10/109,551
CURRENT FILING DATE: 2002-03-28
PRIOR APPLICATION NUMBER: 60/280,549
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                          142 GSDYEDRYY 150
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                                                                                                                                                                                                                                                                                                                              1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: 09/550,374
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELECOMMUNICATION INFORMATION: TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         PRIOR APPLICATION DATA:
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ADDRESSEE: Fish &
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP. NUMBER OF SEQUENCES: 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Prusiner, Stanley B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER: US/09/943,906 FILING DATE: 30-Aug-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FastSEQ Version 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: BOZICEVIC, Karl
REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CITY: Menlo Park
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 253 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TELEFAX: 415-854-0875
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77.8%;
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Pred. No. 0.
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TOPOLOGY: linear

HOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 3:
US-09-943-906-3
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                                                                                                Query Match
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LENGTH: 256
                                                                  Matches
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NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
                                                                                                                                                                                                                                                                   TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 3:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       145 GSDYEDRYY 153
152 GSDYEDRYY 160
                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/09/943,906
FILING DATE: 30-Aug-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/550,374
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: BOZICEVIC, KBI1
REGISTRATION NUMBER: 28,807
REGISTRATION NUMBER: 06510/059001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-5277
TELEPHONE: 415-854-0875
                                                                Local Similarity
nes 7; Conserv
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                               1 GXDXEDRYY 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOETWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Williamson, R. Anthony
Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP
NUMBER OF SEQUENCES: 86
                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT:
                                                                                                                                                                                                                    TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Menlo Park
STATE: CA
                                                                                                                                                                                                 STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 94025
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Prusiner, Stanley B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        93.2%;
77.8%;
                                                                                93.2%;
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                                                                  0.
                                                                              Score 41; DB 1
Pred. No. 0.58;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 41; DB 9
Pred. No. 0.56;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Mismatches
                                                                  Mismatches
                                                                                              DB 10; Length 263;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Length 256;
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Best Local Similarity
Thes 7; Conserve
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US-10-209-194-2
                                                                                                                                                                            RESULT 10
US-09-943-906-4
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.US-10-209-194-2
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                                                                                                                                                                                                                                                                                                                                                                                                 ; ORGANISM: sheep US-09-745-003-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          US-09-745-003-7
                                                                                                     Sequence 4, Application US/09943906 Patent No. US20020150571A1 GENERAL INFORMATION:
APPLICANT: Prusiner, Stanley B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/09/745,003
CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GENERAL INFORMATION:
APPLICANT: LILIFDAHL, MONIKA
APPLICANT: ASPLAND, SIMON ERIC
TITLE OF INVENTION: GENETICALLY MODIFIED COWS HAVING REDUCED
TITLE OF INVENTION: SUSCEPTIBILITY TO MAD COW DISEASE
FILE REFERENCE: BIOBANK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 2, Application US/10209194 Publication No. US20030051264A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 7, Application US/09745003
Patent No. US20020042122A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SOFTWARE:
SEQ ID NO 2
                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION NUMBER: US/10/209,194
CURRENT FILING DATE: 2002-07-29
PRIOR APPLICATION NUMBER: 60/309,222
PRIOR FILING DATE: 2001-07-31
PRIOR APPLICATION NUMBER: 60/67,091
PRIOR APPLICATION NUMBER: 60/67,091
PRIOR FILING DATE: 2002-03-21
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICANT: Bazan, Fernando J
TITLE OF INVENTION: Human Proteins; Related Reagents
FILE REFERENCE: PrP2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQ ID NOS: 15
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TYPE: PRT
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TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 153 GSDYEDRYY 161
                                                                                                                                                                                                                                                      50 GNDYEDRYY 58
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                                                                                                                                                                                                                                                                                       1 GXDXEDRYY 9
                               Williamson, R. Anthony
Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC
NUMBER OF SEQUENCES: 86
                  CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  264
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FastSEQ for Windows Version 4.0
ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                           Conservative
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77.8%;
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Pred. No. 0.55;
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                                                      FOR NATIVE Prp
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STREET: 2200 Sand CITY: Menlo Park

2200 Sand Hill Road

STATE: CA

U.S.A.

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TYPE: amino acids

STRANDEDWESS: single
TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO: 4:
US-09-943-906-4
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                                                                                                                                              ; LENGTH: 256
; TYPE: PRT
; ORGANISM: Ovis aries
US-10-109-551-4
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Best Local Similarity
7; Conserv
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CURRENT FILING DATE: 2002-03-28
PRIOR APPLICATION NUMBER: 60/280,549
PRIOR FILING DATE: 2001-03-30
NUMBER OF SEQ ID NOS: 10
SOFTMARE: PATENTIN VET. 2.1
SEQ ID NO 4
                                                                                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 4, Application US/10109551 Publication No. US20020194635A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                         APPLICANT: DUNNE, PATRICK W.
APPLICANT: PIEDRAHITA, JORGE
TITLE OF INVENTION: TRANSGENIC ANIMALS RESISTANT TO TRANSMISSIBLE
TITLE OF INVENTION: SPONGIFORM ENCEPHALOPATHIES
FILE REFERENCE: TAMK: 307US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
LENGTH: 255 amino acids
145 GNDYEDRYY 153
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                                   1 GXDXEDRYY 9
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible OPERATIER SYSTEM: DOS SOPTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: 09/550,374
FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ZIP: 94025
COMPUTER READABLE FORM:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICATION NUMBER: US/09/943,906 FILING DATE: 30-Aug-2001 CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME: Bozicevic, Karl
REGISTRATION NUMBER: 28,807
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Diskette
                                                                       Conservative
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                                                                                       90.98;
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                                                                                       Pred. No. 0.89;
                                                                      Mismatches
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                                                                                                        Length 256;
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RESULT 14
US-10-109-551-10
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TYPE: PRT
; ORGANISM: Odocoileus hemionus hemionus
US-10-109-551-8
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Best Local Similarity
Thes 7; Conserv
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; SEQ ID NO 6
; LENGTH: 256
; TYPE: PRT
; ORGANISM: Odocoileus virginianus
US-10-109-551-6
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 13
US-10-109-551-8
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US-10-109-551-6
Sequence 10, Application US/10109551
Publication No. US20020194635A1
GENERAL INFORMATION:
APPLICANT: DUNNE, PATRICK W.
APPLICANT: PIEDRAHITA, JORGE
TITLE OF INVENTION: TRANSGENIC ANIMALS RESISTANT TO TRANSMISSIBLE
TITLE OF INVENTION: SPONGIFORM ENCEPHALOPATHIES
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 8, Application US/10109551 Publication No. US20020194635A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. SEQ ID NO 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Sequence 6, Application US/10109551
Publication No. US20020194635A1
GENERAL INFORMATION:
APPLICANT: DUNNE, PATRICK W.
APPLICANT: PIEDRAHITA, JORGE
                                                                                                                                                                                                                                                                                                                         Best Local Similarity Matches 7; Conserv
                                                                                                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CURRENT APPLICATION NUMBER: US/10/109,551
CURRENT FILING DATE: 2002-03-28
PRIOR APPLICATION NUMBER: 60/280,549
PRIOR FILING DATE: 2001-03-30
NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: DUNNE, PATRICK W.
APPLICANT: PIEDRAHITA, JORGE
TITLE OF INVENTION: TRANSGERIC ANIMALS RESISTANT TO TRANSMISSIBLE
TITLE OF INVENTION: SPONGIFORM ENCEPHALOPATHIES
FILE REFERENCE: TAMK:207US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIOR APPLICATION NUMBER: 60/280,549 PRIOR FILING DATE: 2001-03-30 NUMBER OF SEQ ID NOS: 10
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/109,551
CURRENT FILING DATE: 2002-03-28
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                                                                                                                                                                                                                                          145 GNDYEDRYY 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    145 GNDYEDRYY 153
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Pred. No. 0.89;
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                                                                                                                                                                                                                                                                                                                                                                   Length 256;
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Sequence 11, Application US/09745003

Patent NO. US2020042122A1

GENERAL INFORMATION:
APPLICANT: Bazan, Fernando J

TITLE OF INVENTION: Human Proteins; Related Reagents
FILE REFERENCE: PTP2

CURRENT APPLICATION NUMBER: US/09/745,003

CURRENT FILING DATE: 2000-12-20

NUMBER OF SEQ ID NOS: 13

SOFTWARE: Patentin Ver. 2.0

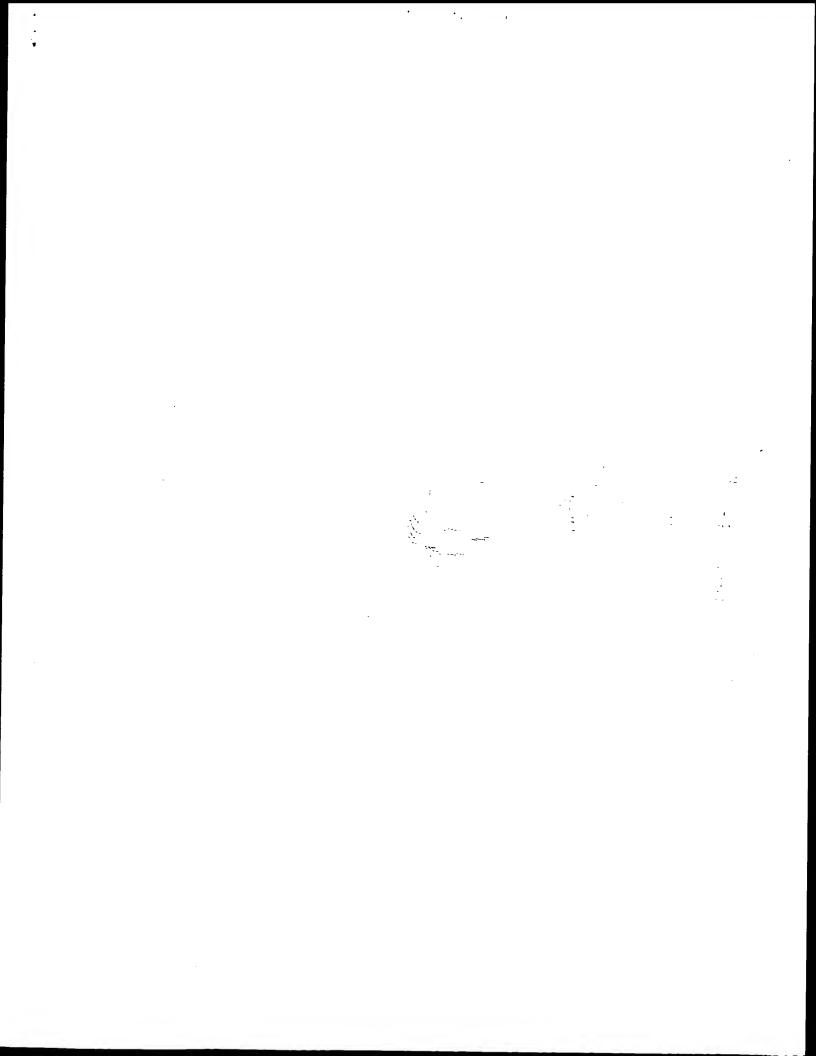
SEQ ID NO 11

LENGTH: 163

TYPE: PRT

ORGANISM: Hamster sp.

US-09-745-003-11
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILE REFERENCE: TAMK:207US
CURRENT APPLICATION NUMBER: US/10/109,551
CURRENT FILING DATE: 2002-03-28
PRIOR APPLICATION UTWEER: 60/280,549
PRIOR FILING DATE: 2001-03-30
NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 10
LENGTH: 256
TYPE: PRT
ORGANISM: Cervus elaphus
US-10-109-551-10
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US-09-745-003-11
Search completed: March 24, 2003, 17:25:04 Job time: 9.0625 secs
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Best Local Si
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Best Local Similarity 77.8%;
Matches 7; Conservative
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| | ||||
145 GNDYEDRYY 153
                                                                    1 GXDXEDRYY 9
| | ||||
51 GNDWEDRYY 59
                                                                                                                                     y Match 88.6%;
Local Similarity 77.8%;
hes 7; Conservative
                                                                                                                                          Score 39; DB 10; Length 163; Pred. No. 0.88; 0; Mismatches 2; Indels
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Pred. No. 0.89;
0; Mismatches
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                                                                                                                                                0; Gaps
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GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.

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Minimum DB :
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence:
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seq length: 2000000000
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SUMMARIES

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51	51	51	51	51	51	51	51	51	51	Score
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AAR38010 standard; protein; 27 ₿

Prion protein region C #4.

14-OCT-1993 (first entry)

Antigen; prion; protein; region; frame shift; repeat; mutation; PrPc; FSa; FSb; subfragment; antibody; treatment; spongiform encephalopathy; human; sheep; cattle; cellular binding; aggregation; mammal; scrapie; immune system; PrPsc; ratio-inverso peptide; enzymatic degradation; resistance.

Synthetic.

W09311155-A.

10-JUN-1993.

03-DEC-1992; 92WO-GB02246

03-DEC-1991; 10-JUL-1992; 91GB-0025747. 92GB-0014663.

(PROT-) PROTEUS MOLECULAR DESIGN LTD.

Fishleigh RV, Mee RP,

WPI; 1993-196994/24.

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   The sequences given in AAR38007-11 from an antigenic site, region C,
                                                Claim 15; Page 67; 82pp; English.
                                                                                                New polypeptide(s) contg. antigenic site of prion protein - useful for treatment and diagnosis of mammalian encephalopathies
                                                                                                                                                                                                                                                                                                                                                                                                                                               Antigen; prion; protein; region; frame shift; repeat; mutation; PrPc; FSa; FSb; subfragment; antibody; treatment; spongiform encephalopathy; human; sheep; cattle; cellular binding; aggregation; mammal; scrapie; immune system; PrPsc; ratio-inverso peptide; enzymatic degradation;
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10-JUL-1992;
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92GB-0014663.
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69.2%;
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10-JUL-1992;
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92GB-0014663.
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69.2%;
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Pred. No. 0.0027;
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Synthetic

Misc-difference

Location/Qualifiers

/note= "One or more residues or may be absent"

Misc-difference

Prion protein region C #1.

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RESULT 6
AAR38007
ID AAR3
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Best Local :
                                                                                                                                                                                               The sequences given in AAR38012-15 represent polypeptide subfragments derived from an antigenic site, region C, of a prion protein. Prion proteins comprise six regions of interest (A-F), and two related frame shift peptides sequences caused by a repeating section in region E having a nucleic acid coding sequence frame shift mutation of +1 (FSa) or -1 (FSb). These subfragments or the full length peptides (see AAR38006-11), and antibodies raised against these, may be used to treat or prevent spongiform encephalopathy in humans, sheep or cattle. They can be used to block cellular binding and aggregation of prion proteins and to stimulate the mammalian immune system. These peptides may be used to distinguish between the normal form of prion protein (PrPc) and the scrapic-associated form (PrPsc). These peptides
   14-OCT-1993
                                                                                                                                                                                                                                                                                                                                         New polypeptide(s) contg. antigenic site of prion protein - useful for treatment and diagnosis of mammalian encephalopathies e.g. Creutzfeid-Jacob disease and kuru
                      AAR38007;
                                       AAR38007 standard; protein; 31 AA
                                                                                                                                                                Sequence
                                                                                                                                                                              protein (PrPc) and the scrapic associated form (PrPsc). These peptimay include rare or synthetic amino acids or a ratio inverso peptide modification to improve resistance to enzymatic degradation.
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10-JUL-1992;
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9; Conserv
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Query Match
Best Local Similarity
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The sequences given in AAR38007-11 represent polypeptides derived from an antigenic site, region C, of a prion protein. Prion proteins somprise six regions of interest (A-F), and two related frame shift peptides sequences caused by a repeating section in CC region E having a nucleic acid coding sequence frame shift mutation CC of +1 (FSa) or -1 (FSb). These peptides and subfragments of these CC (see AAR38012-15), and antibodies raised against these, may be used to treat or prevent spongiform encephalopathy in humans, sheep or cattle. They can be used to block cellular binding and aggregation of prion CC proteins and to stimulate the mammalian immune system. These peptides may be used to distinguish between the normal form of prion protein CC (PrPc) and the scrapic-associated form (PrPsc). These peptides may CC include rare or synthetic amino acids or a ratio-inverso peptide composition to improve resistance to enzymatic degradation.
Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 14; Page 66; 82pp; English.
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10-JUL-1992;
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The sequences given in AAR38007-11 represent polypeptides derived from an antigenic site, region C, of a prion protein. Prion proteins comprise six regions of interest (A-F), and two related frame shift peptides sequences caused by a repeating section in region E having a nucleic acid coding sequence frame shift mutati
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    eating section in frame shift mutation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            may be
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Best Local 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         of +1 (FSa) or -1 (FSb). These peptides and subfragments of these (see AAR38012-15), and antibodies raised against these, may be used to treat or prevent spongiform encephalopathy in humans, sheep or cattle. They can be used to block cellular binding and aggregation of prion proteins and to stimulate the mammalian immune system. These peptides may be used to distinguish between the normal form of prion protein (PrPc) and the scrapic-associated form (PrPsc). These peptides may include rare or synthetic amino acids or a ratio-inverso peptide modification to improve resistance to enzymatic degradation.
                                                                                                                                                                                                                                                                                                                                                                                                  /note-
Misc-difference 29
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Antigen; prion; protein; region; frame shift; repeat; mutation; PrPc; FSa; FSb; subfragment; antibody; treatment; spongiform encephalopathy; human; sheep; cattle; cellular binding; aggregation; mammal; scraple; immune system; PrPsc; ratio-inverso peptide; enzymatic degradation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Synthetic.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       AAR38009 standard;
           Fishleigh RV,
                                                                                                 03-DEC-1991;
10-JUL-1992;
                                                                                                                                                                                                                                                                                                                                  Misc-difference
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                                                                                                                                                                       03-DEC-1992;
                                                    (PROT-) PROTEUS
                                                                                                                                                                                                               10-JUN-1993.
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9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                           /note=
           Mee RP,
                                                                                                    91GB-0025747.
92GB-0014663.
                                                                                                                                                                       92WO-GB02246
                                                         MOLECULAR DESIGN LTD.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Location/Qualifiers
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69.2%;
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           Robson B;
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Pred. No. 0
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                                                                                                                                                                                                                                                                                                                                                                                                                                               absent"
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0.003;
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RESULT 9
AAW35545
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      A non-dendritic peptide carrier (A) has been developed which is coupled through a linker to a solid phase, forming a complex of (A)-solid phase Where (A) comprises 10.50 amino acids capable of forming a secondary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            from an antigenic site, region C, of a prion protein. Prion proteins comprise six regions of interest (A-F), and two related frame shift peptides sequences caused by a repeating section in region E having a nucleic acid coding sequence frame shift mutation of +1 (FSb) or -1 (FSb). These peptides and subfragments of these (see AAR38012-15), and antibodies raised against these, may be used to treat or prevent spongiform encephalopathy in humans, sheep or cattle. They can be used to block cellular binding and aggregation of prion proteins and to stimulate the manmalian immune system. These peptides may be used to distinguish between the normal form of prion protein (PrPc) and the scrapic-associated form (PrPsc). These peptides may include rare or synthetic amino acids or a ratio-inverso peptide
                                                                    Example 31; Page
                                                                                                         Non-dendritic peptide carrier linked to a solid phase - useful as a diagnostic agent and as a scaffold for production of chemical
                                                                                                                                                             WPI; 1997-512645/47.
                                                                                                                                                                                                                            (PEPR-) PEPRESEARCH AS.
                                                                                                                                                                                                                                                          03-APR-1996;
                                                                                                                                                                                                                                                                                         03-APR-1997;
                                                                                                                                                                                                                                                                                                                                                   WO9738011-A1
                                                                                                                                                                                                                                                                                                                                                                                                         T-cell stimulatory peptide; immunogen; non-dendritic; carrier; tumour; scaffold; inhibition; metastasis; wound healing; solid phase.
                                                                                                                                                                                                                                                                                                                      16-OCT-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        22-APR-1998 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   AAW35545 standard; peptide; 33
                                                                                                                                                                                                                                                                                                                                                                                                                                                         Immunization DNDPC SEQ ID NO:89 from WO9738011.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   The sequences given in AAR38007-11 represent polypeptides derived
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 14; Page 67; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    e.g. Creutzfeld-Jacob disease and kuru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New polypeptide(s) contg. antigenic site of prion protein useful for treatment and diagnosis of mammalian encephalopathies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LOCAL
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9; Conser
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                                                                                                                                                                                            Jakobsen PH;
                                                                                                                                                                                                                                                        96DK-0000398
                                                                                                                                                                                                                                                                                     97WO-DE00146.
                                                                  156;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87.9%;
69.2%;
                                                                262pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 51; DB 14
Pred. No. 0.003;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      3
   acids capable of
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forming a secondary
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cc structure in a benign buffer after liberation from the solid phase, and cc further the (A)-solid phase complex comprises an immunogenic substance and/or an immune mediator coupled on (A). The present sequence cc represents a peptide used in an example from the present invention. An cc (A)-solid phase complex can be used as a scaffold for the production of cc chemical derivatives, characterised by covalently attaching molecules at attachment points. Alternatively (A) is used as a scaffold peptide for the incorporation into an immunostimulating Complex (Iscom) resulting an cc (A)-iscom complex which is used for the chemical coupling of antigenic substances in an aqueous solution by conjugation. (A) derivatised with cone or more peptides having fibronectin. Jaminin- or vitronectin-like binding activities can be used for the promotion of cell-attachment to classify surfaces, in particular to inhibit tumour growth and metastasis, and for promotion of specifically-binding aptamers or as a diagnostic agent. Such diagnostic-(A) molecules could be used to detect molecules conference into a inferious and an anti-immune or concerns disease.
                                                    infectious, autoimmune or cancerous disease.
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밁 Matches Query Match 19 QVYYRPVDQYSNQ Local Similarity 1 QVYYXPXDXYXNQ 13 9; Conservative 87.9%; 69.2%; 0; Score 51; Pred. No. Mismatches DB 18; Length 33; 0 Gaps

0

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Sequence

33 AA;

Prion; PrP; guanidine thiocyanate; gdnSCN; TSE; BSE; transmissible spongiform encephalopathy; antibody; bovine spongiform encephalopathy; sheep; cattle; human; mouse; 18-DEC-2000 (first entry) Homo sapiens. Human prion protein peptide homologous to ovine sequence 145-177. AAB15054 standard; Peptide; hamster; rabbit.

WO200048003-A1

17-AUG-2000.

09-FEB-2000; 2000WO-NL00079.

11-FEB-1999; 99EP-0200391.

(DIEN-) STICHTING DIENST LANDBOUWKUNDIG ONDERZOE.

Schreuder BEC, Garssen GJ, Jacobs JG, L C, Bossers A; Langeveld JPM, Smits MA, Van Keulen LJM;

WPI; 2000-506099/45.

Use of guanidine thiocyanate for reducing risk of false-positive results in testing mammalian sample for aberrant prion protein, useful for detection of transmissible spongiform encephalopathies -

Disclosure; Fig 2; 49pp; English.

The present invention relates to a method for reducing the risk of scoring a false positive test result in testing a sample for aberran prion protein. The method involves the use of guanidine thiocyanate (gdnSCN) or its functional equivalent. This test is highly useful for spongiform encephalopathies for aberrant

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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
                                                                                                      The present invention relates to a method for reducing the risk of scoring a false positive test result in testing a sample for abbrrant prion protein. The method involves the use of guandine thiocyanate (gdnSCN) or its functional equivalent. This test is highly useful for testing for transmissible spongiform encephalopathies (TSEs) such as BSE (bovine spongiform encephalopathy). The method allows a faster, simpler and more reliable method for monitoring cattle and sheep for the presence of abarrant prion protein before it reaches the human and animal food chain. In the invention antipeptide antibodies were raised against sheep prion protein peptides. The present sequence is the rabbit prion protein sequence homologous to the sheep peptide
                                                                                                                                                                                                                                                                                                                                                                         Use of guanidine thiocyanate for reducing risk of false-positive results in testing mammalian sample for aberrant prion protein, useful for detection of transmissible spongiform encephalopathies -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            BSE (bovine spongiform encephalopathy). The method allows a faster, simpler and more reliable method for monitoring cattle and sheep for the presence of aberrant prion protein before it reaches the human and animal food chain. In the invention antipeptide antibodies were raised against sheep prion protein peptides. The present sequence is the human prion protein sequence homologous to the sheep peptide
                                                         Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              11-FEB-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  09-FEB-2000; 2000WO-NL00079
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Oryctolagus cuniculus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prion; PrP; guanidine thiocyanate; gdnSCN; TSE; BSE; transmissible spongiform encephalopathy; antibody; bovine spongiform encephalopathy; sheep; cattle; human.
                                                                                                                                                                                                                                                                                                                                           Disclosure; Fig 2; 49pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-506099/45.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Garssen GJ,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            AAB15055 standard; Peptide; 33
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (DIEN-) STICHTING DIENST LANDBOUWKUNDIG ONDERZOE.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Rabbit prion protein peptide homologous to ovine sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               18-DEC-2000 (first entry)
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                                                         33 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Jacobs JG,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Bossers
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69.2%;
 87.9%;
69.2%;
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Pred. No.
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Pred.
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51;
No.
 DB 21;
0.0032;
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                 Length 33;
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RESULT 13 AAB15058

AAB15058 standard; Peptide; 33 AA.

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RESULT 12
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                                                                                                                                          Sequence
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WO200048003-A1
                                                                                Prion; PrP; guanidine thiocyanate; gdnSCN; TSE; BSE; transmissible spongiform encephalopathy; antibody;
                                                                                                                                    Sheep prion protein peptide
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                                                                  bovine spongiform encephalopathy; sheep; cattle; human
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                                                                                                                                                                                                                                                                                      Prion protein; PrP; alpha helical domain; screening; inhibition; binding; scrapie; bovine spongiform encephalopathy; BSE; CJD; Creutzfeldt-Jakob disease; kuru; GSS; FFI; fatal familial insomn
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
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                              Cohen FE, Kaneko K,
                                                                                                  02-NOV-1995;
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                                                                                                                                                                                                        WO9716728-A1
                                                                                                                                                                                                                                                                          Gerstmann-Straussler-Scheinker disease;
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                                                                (REGC ) UNIV CALIFORNIA.
                                                                                                                                     28-OCT-1996;
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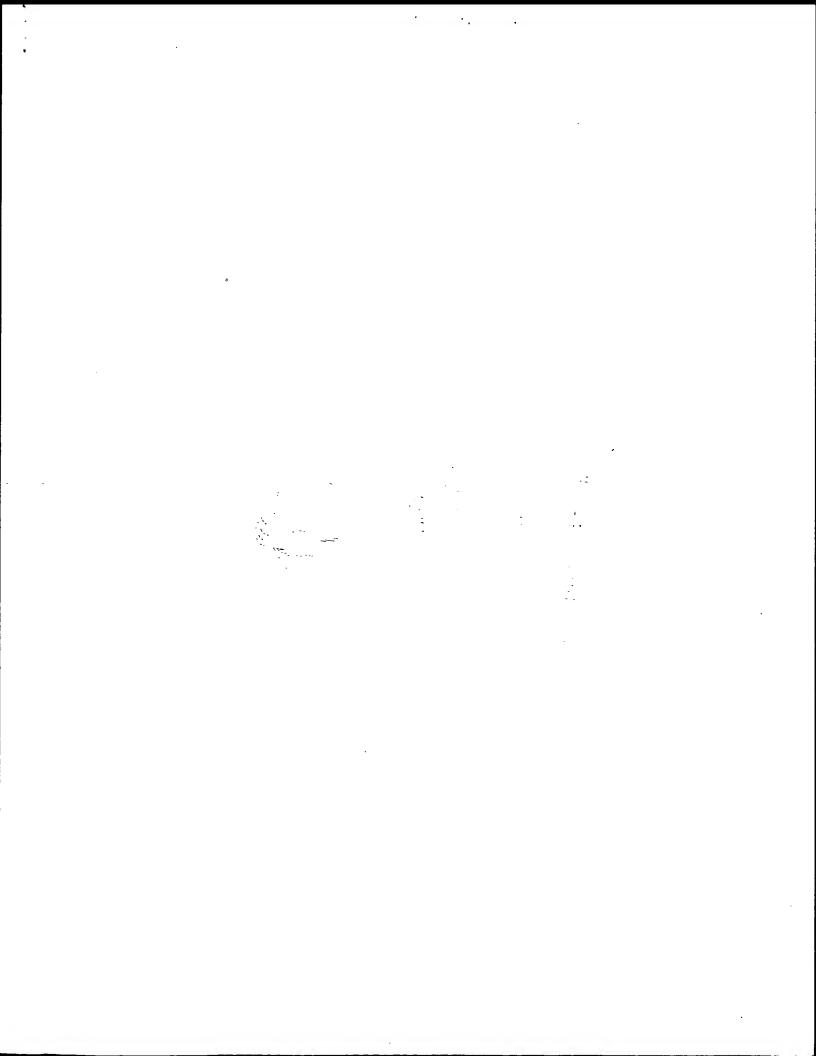
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Prion proteins (PrPs) having at least one alpha-helical domain - Pr used in assays for screening compounds able to inhibit or decrease Pr the binding of PrP peptide(s) to cellular prion proteins or peptide(s)

XX Claim 11; Page 7-38; 50pp: English.

XX The present sequence represents a prion protein (PrP) peptide. CC PrP has an ability to induce a conformational change in cellular CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC prion protein (PrP-c). Methods, for screening compounds which CC protein protein (PrP-c). Methods for screening compounds which CC protein protein (PrP-c). Methods for screening compounds which CC protein protein (PrP-c). Methods, for screening compounds which CC protein protein (PrP-c) Methods, for screening compounds which CC protein protein (PrP-c) pepti
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Result No.

Score 51;

DB 2;

Length 239;

RESULT 2 S53633 major prik C; Species C; Date: 2 Anccessi A, Mol. B. A; Title: 1 A; Residues A; Cross-ro A; Residues A; Referend A; Referend A; Recessi A; Molecul A; Residues A; Cross-ro C; Superfan C; Keyword C; Keyword C; Status C	RESULT 1 A53892 prion-rela C;Species C;Date: 0; C;Accessive A;Tilao, Y Lab. Inves A;Tila: (A;Accessive A;Accessive	33 33 33 33 33 33 33 33 33 33 33 33 33
	Palated protein - rat (fragment) ss: Rattus norvegicus (Norway rat) 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 13-Aug-1999 ion: A53892 Y.C.; Tokes, Z.; Lim, E.; Lackey, A.; Woo, C.H.; Button, J.D.; Clawson, G. rest. 57, 370-374, 1987 rect. 57, 370-374, 1987 rect. 57, 370-374, 1987 rect. 57, 370-374, 1987 prion related protein cDNA. noce number: A53892; MUID:88037055; PMID:2889848 sion: A53892 ipreliminary ile type: mRNA les: 1-226 <lia> references: GB:M20313; NID:g206391; PIDN:AAA41947.1; PID:g206392 family: major prion protein Match Ocal Similarity 69.2%; Score 51; DB 2; Length 226; ocal Similarity 69.2%; Pred. No. 0.005; ss 9; Conservative 0; Mismatches 4; Indels 0; Gaps 0; 1 QVYYXPXDXYXNQ 13 1 1 </lia>	30 50 86.2 232 2 S71041 major prion protei 31 50 86.2 252 2 S53634 major prion protei 32 50 86.2 254 1 UJHYIH major prion protein - Ch 33 50 86.2 254 2 A34759 prion protein - Ch 34 50 86.2 257 2 A23545 prion protein - Go 35 46 79.3 253 2 S53635 prion protein protei 37 46 79.3 253 2 S53635 prion protein protei 39 46 79.3 253 2 S53635 prion protein protei 39 40 2 S76066 prion protein - Si 40 37 63.8 295 2 B82674 prion protein protein 41 37 63.8 458 2 T25115 prion protein protein 42 36 62.1 146 2 E69773 protein protein protein 43 36 62.1 396 2 S42225 prion protein pr
e e	>	

Best Local Matches

Similarity 69.: 9; Conservative

69.2%;

Pred. No. 0.0053;

Mismatches

4;

Indels

0

Gaps

0;

밁

153

QVYYRPVDQYSNQ 165

0;

μ

QVYYXPXDXYXNQ 13

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A;Status: nucleic acid sequence not shown
A;Molecule type: DNA
A;Residues: 1-203, 'R', 205-240 <SCW>
A;Cross references: EMBL:U08312
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             R:Schatzl, H.M. submitted to the EMBL Data Library, April 1994 A:Reference number: S71041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       A;Title: Prion protein gene variation among primates.
A;Reference number: $53614; MUID:95139066; PMID:7837269
A;Recession: $53621
A;Status: nucleic acid sequence not shown
A;Nolecule type: DNA
A;Nolecule type: DNA
A;Residues: 1-203; Kr. 205-240 <SCW>
A;Cross-references: EMBL:008303
A;Cross-references: EMBL:008303
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
                       á
                                                                                                                                                                                                                                                                                                    A;Cross-references: EMBL:U08312; NID:g475585; PIDN:AAC50100.1; PID:g475586 F;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B. J. Mol. Biol. 245, 362-374, 1995
J. Mol. Biol. 245, 362-374, 1995
A;Title: Prion protein gene variation among primates.
A;Reference number: S53614; MUID:95139066; PMID:7837269
A;Accession: S53632
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submitted to the EMBL Data Library, April 1994
A:Reference number: S71041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           major prion protein - mandrill (fragment)
c;Species: Papio sphinx, Mandrillus sphinx (mandrill)
C;Date: 27-Oct-1996 #sequence_revision 14-Feb-1997 #t.
C;Accession: S71056; S53621
                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Molecule type: DNA
A; Residues: 1-241 <SCH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Accession:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   major prion protein - Callicebus moloch (fragment)
C;Species: Callicebus moloch
C;Date: 27-Oct-1996 #sequence_revision 07-Feb-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           S71048
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          A; Molecule type: DNA
A; Residues: 1-241 <SCH>
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Best Local
                                                                         Matches
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                                                                       Local Similarity
mes 9; Conserv
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                          QVYYXPXDXYXNQ 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                S71048; S53632
                                                                       Conservative
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69.2%;
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69.2%;
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                                                                                       Score 51; DB 2; Length 241; Pred. No. 0.0053;
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                                                                    Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
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                                                                    Indels
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161848
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A;Residues: 8-10,'L',12-202,'/R',204-239 <SCW>
A;Cross references: EMBL:U08892
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scraple; transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        R;Schatzl, H.M.
submitted to the EMBL Data Library,
""Deference number: S71041
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A;Accession: S53628
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A; Residues: 1-245 <SCH>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Species: Cercopithecus dianae
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 13-Aug-1999
C;Accession: S71045; S53628
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            major prion protein - Cercopithecus dianae
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                                                                                                                                                                                                                                                                                                                                                                                     A; Status: nucleic acid sequence not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A; Cross-references: EMBL: U08292; NID: g474342; PIDN: AAC50081.1; PID: g474343
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                                                                                                                                                                               Matches
                                                                                                                                                                                                                             Query Match
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                                                                                       152 QVYYRPVDQYSNQ 164
                                                                                                                                  1 QVYYXPXDXYXNQ 13
                                                                                                                                                                                  9;
                                                                                                                                                                                  Conservative
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87.9%;

Score 51; DB 2; Pred. No. 0.0054;

Length 245;

00

Mismatches

0

0

Taylor, L.; Cohen, F.E.; Prusiner, S.B.

PMID: 7837269

April 1994

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A; Molecule type: DNA
A; Residues: 1-10,'v',12-202,'E',204-245 <SCW>
A; Residues: 1-10,'v',12-202,'E',204-245 <SCW>
A; Cross references: EMBL:U08891; NID:g474340; PIDN:AAC50080.1; PID:g474341
C; Superfamily: major prion protein
C; Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                  submitted to the EMBL Data Library, April 1994 A; Reference number: S71041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             A;Title: Prion protein gene variation among primates. A;Reference number: S53614; MUID:95139066; PMID:7837269 A;Accession: S53627
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              C;Species: Cercopithecus aethiops (green monkey, grivet)
C;Date: 28-Oct-1996 #sequence_revision 07-Feb-1997 #text_change 13-Aug-1999
C;Accession: S53627; S71043
R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.
J. Mol. Biol. 245, 362-374, 1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        A; Molecule type: DNA
A; Residues: 1-245 <SCH>
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152 QVYYRPVDQYSNQ 164
                                                                                                                      Local Similarity
les 9; Conser
                                                        1 QVYYXPXDXYXNQ 13
                                                                                                                      Conservative
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                                                                                                                                            87.9%;
69.2%;
                                                                                                                   0,:
                                                                                                                                            Score 51; DB 2; Length 245; Pred. No. 0.0054;
                                                                                                                   Mismatches
                                                                                                                0
                                                                                                                Gaps
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.A;Molecule type: DNA
A;Residues: 1-209,'E',211-252 <SCW>
A;Residues: 1-209,'E',211-252 <SCW>
A;Cross-references: EMB::U08295; NID:g474348; PIDN:AAC50084.1; PID:g474349
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane
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A;Title: Infectious amyloid precursor gene sequences in primates used for experimental A;Reference number: 136907; MUID:95083661; PMID:7991600
A;Accession: 161848
A; Molecule type: DNA
A; Residues: 1-252 <LDS-
A; Cross references: GB:UZ8334; NID:gl490412; PIDN:AAC48697.1;
C; Comment: This protein is a cellular protein, it is involved
                                                                                                       A;Title: Characterization of a prion protein (PrP) gene from rabbit; a species with apportance number: JC6175; MUID:97183665; PMID:9031631
A;Accession: JC6175
                                                                                                                                                                                                                                  prion protein - rabbit
C;Species: Oryctolagus cuniculus (domestic rabbit)
C;Date: 11-Apr-1997 #sequence_revision 09-May-1997 #text_change 13-Aug-1999
C;Accession: JC6175
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     A;Title: Prion protein gene variation among primates. A;Reference number: S53614; MUID:95139066; PMID:7837269 A;Accession: S53631
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C;Species: Cebus apella (brown capuchin, black-capped capuchin) C;Date: 28-Oct-1996 *sequence_revision 07-Feb-1997 *text_change C;Accession: S53631; S71044
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C;Dato: 31-May-1996 #sequence_revision 31-May-1996 #text_change 13-Aug-1999
                                                                                                                                                                               R; Loftus, B.; Rogers, M. Gene 184, 215-219, 1997
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A; Accession: S71044
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J. Mol. Biol. 245, 362-374, 1995
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A; Residues: 1-252 < RES>
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                                                                                                                                                                                                                                                                                                                                                                    RESULT 9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     major prion
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Status: preliminary; translated from GB/EMBL/DDBJ
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Local Similarity
les 9; Conserv
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les 9; Conserv
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69.2%;
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69.28;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Score 51; DB 2;
Pred. No. 0.0056;
0; Mismatches
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Pred. No. 0.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 252;
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  PID:g1490413
in the neuro
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A;Status: preliminary; translated from GB/EMBL/DDBJ
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A; Residues: 1-80,89-253 <PUC>
A; Cross-references: GB:X83416; NID:g747846; PIDN:CAA58442.1; PID:g747847;
A; Note: the deletion may be a polymorphism; the alternative deletion of (R;Liao, Y.C.J.; Lebo, R.V.; Clawson, G.A.; Smuckler, E.A. Science 233, 364-367, 1986
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                R; Puckett, C.; Concannon, P.; Casey, C.; Hood, L. Am. J. Hum. Genet. 49, 320-329, 1991
A; Title: Genomic structure of the human prion protein A; Reference number: A40372; MUID:91328137; PMID:167824
A; Accession: A40372
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     C;Species: Homo sapiens (man)
C;Date: 25-Oct-1987 #sequence_revision 12-Apr-1996 #text_change 16-Jun-2000
C;Accession: A24173; A40372; A05017; S14078; I54322; I68597; I58135; I59184.
R;Kretzschmar, H.A.; Stowring, L.E.; Westaway, D.; Stubblebine, W.H.; Prusin DNA 5, 315-324, 1986
A;Cross-references: GB:M81930; NID:g190519; PIDN:AAB59443.1; R;Brown, P.; Goldfarb, L.G.; McCombie, W.R.; Nieto, A.; Squil Neurology 42, 422-427, 1992
A;Title: Atypical Creutzfeldt-Jakob disease in an American fa A;Reference number: IS8135; MUID:92140671; PMID:1736177
A;Accession: IS8135
                                                                                                                                                                                                                                                                                                                                                                                                                                  A;Molecule type: protein
A;Residues: 58-72,'x',74-76,'xx',79,'xxx',83-86;111-128,'V',130-150
A;Residues: 58-72,'x',74-76,'xx',79,'xxx',83-86;111-128,'V',130-150
Hum. Mol. Genet. 1, 443-444, 1992
A;Title: Deletion in the prion protein gene in a demented patient.
A;Reference number: 154322; MUID:93250789; PMID:1363802
A;Accession: 154322
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Molecule type: mRNA
A;Residues: 8-117,119-253 <LIA>
A;Residues: 8-117,119-253 <LIA>
A;Cross-references: GB:D00015; NID:g220015; PIDN:BAA00011.1; PID:g220016; GB:M13667;
R;Tagliavin1, F.; Prelli, F.; Ghiso, J.; Bugian1, O.; Serban, D.; Prusiner, S.B.; Fa
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A;Cross-references: GB:M13899; NID:g190467; PIDN:AAA60182.1; PID:g190468 R;Puckett, C.; Concannon, P.; Casey, C.; Hood, L. Am. J. Hum. Genet. 49, 320-329, 1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    major prion protein precursor - human
N;Alternate names: 11K amyloid protein; 27-30K sialoglycoprotein; PrP 27-30; PrP 33-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    C; Superfamily: major prion protein
C; Keywords: disulfide bond; prion
                                                                                                                                                                                  A; Residues: 8-240 < RE3>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Title: Amyloid protein of Gerstmann-Straeussler-Scheinker disease (Indiana kindred) A;Reference number: S14078; MUID:91160504; PMID:1672107
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A; Reference number: A05017; MUID: 86261778; PMID: 3014653 A; Accession: A05017
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A; Residues: 1-253 <KRE>
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                                                                                                                                                                                                                                                                                    A;Cross-references: GB:M81929; NID:g190517; PIDN:AAB59442.1; PID:g190518
A;Accession: I68597
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                                                                                                                                                                                                                      A; Molecule type: DNA
                                                                                                                                                                                                                                                   A;Status: translated from GB/EMBL/DDBJ
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A;Residues: 9-83,92-240 <RES>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity hes 9; Conserv
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Pred. No. 0.
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Squillacote, D.;
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C;Species: Macaca mulatta (rhesus macaque)
C;Date: 24-May-1996 #sequence_revision 24-May-1996 #text_change 13-Aug-1999
C;Accession: I84423; S53622; S71054
C;Accession: I84423; S53622; S71054
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A; Residues: 51-91, PHGGGWGQPHGGGWGQPHGGGWGQPHGGGWGQPHGGGWGQPHGGG, <RE2>
A; Cross-references: GB:S80539; NID:g244698; PIDN:AAB21334.1; PID:g244699
A; Cross-references: GB:S80539; NID:g244698; PIDN:AAB21334.1; PID:g244699
A; Goldfarb, L.G.; Brown, P.; McCombie, W.R.; Goldgaber, D.; Swergold, G.D.; Wills, P.R
Proc. Natl. Acad. Sci. U.S.A. 88, 10926-10930, 1991
A; Title: Transmissible familial Creutzfeldt-Jakob disease associated with five, seven, A; Reference number: I59184; MUID:92073400; PMID:1683708
A; Accession: I59184
                                                                                                                                     major prion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A;Title: Infectious amyloid precursor gene sequences in A;Reference number: I36907; MUID:95083661; PMID:7991600 A;Accession: I37032
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Date: 31-May-1996 #sequence_revision 31-May-1996 #text_change 13-Aug-1999 C;Accession: I37032
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A;Note: this gene is associated with Creutzfeld-Jakob disease (CJD), Gerstmann-Strausler
C;Superfamily: major prion protein
C;Keywords: amyloid; blocked carboxyl end; brain; glycoprotein; lipoprotein; phosphatidy
F;1-22/Domain: signal sequence #status predicted <SIG>
F;23-230/Product: major prion protein #status predicted <MAT>
F;54-92/Region: 8-residue repeats (P-H-G-G-G-W-G-O)
                              R; Cervenakova,
                                                                                                                                                                                         RESULT 12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                C; Superfamily: major prion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           R;Cervenakova, L.; Brown, P.; Goldfarb, L.G.; Nagle, J.; Pettrone, Proc. Natl. Acad. Sci. U.S.A. 91, 12159-12162, 1994
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F;181,197/Binding site: carbohydrate (Asn) (covalent)
F;230/Modified site: GPI-anchor ethanolamine amidated
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A;Map position: 20pter-20p12
A;Introns: #status absent
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A; Residues: 60-67 <GOL>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Molecule type: DNA
Residues: 1-253 <RES>
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                                                                                                                                                                                                                                                                                                                                                                              Matches
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Best Local
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                                                                                                                                                                                                                                                                      160
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                                                                                                                                                                                                                                                                                                                                                                                                      Local
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                                                                                                                                                                                                                                                                      QVYYRPMDQYSNQ 172
  va, L.; Brown, P.; Goldfarb, L.G.; Nagle, Acad. Sci. U.S.A. 91, 12159-12162, 1994
                                                                                                                                protein precursor - rhesus macaque
                                                                                                                                                                                                                                                                                                                                                                              Similarity
9; Conserv
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9; Conserva
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69.2%;
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Pred. No.
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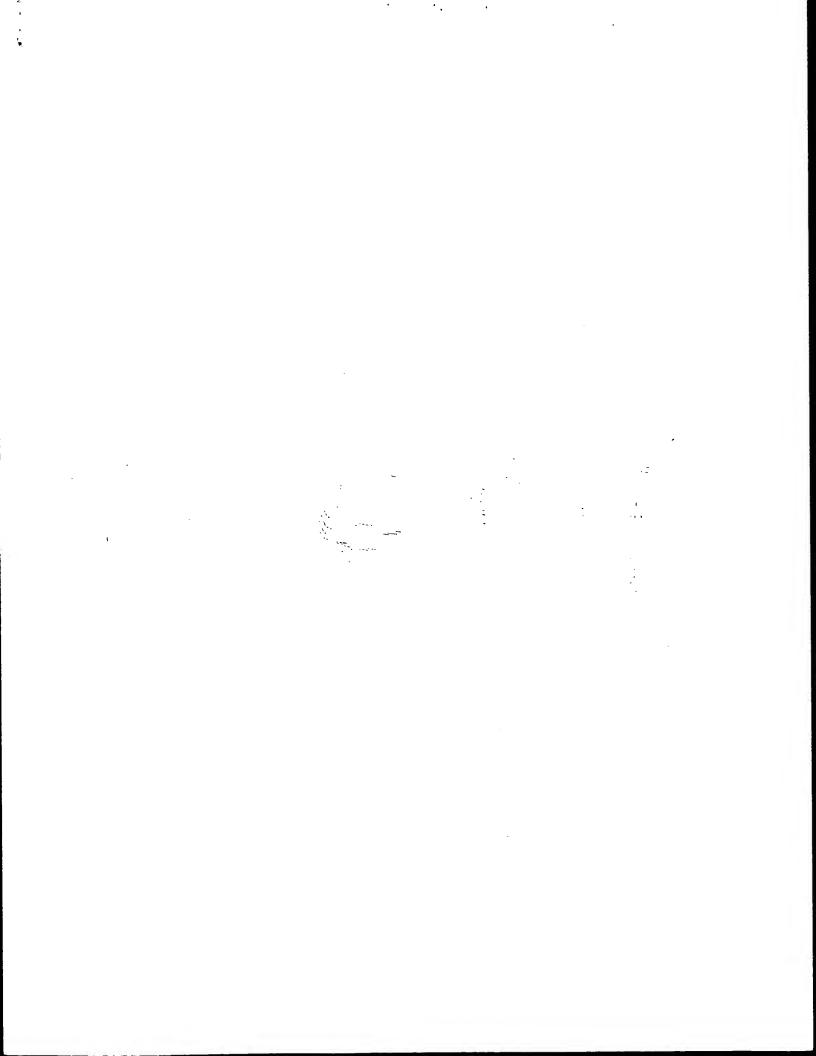
Presbytis

Presbytis

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A; Molecule type: DNA
A; Residues: 1-210, 'E', 212-253 <SCW>
A; Cross references: EMBL:U08297; NID:g474352; PIDN:AAC50086.1; PID:g474353
C; Superfamily: major prion protein protein; lipoprotein; prion; scrapie; traic; Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:U08307; NID:g474372; PIDN:AAC50095.1; PID:g474373 C;Superfamily: major prion protein C;Keywords: amyloid; brain; glycoprotein; llpoprotein; prion; scrapie; tra
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                                                                                                                                                                                                                                                                                                                                           submitted to the EMBL Data A; Reference number: S71041
                                                                                                                                                                                                                                                                                                                                                                                                                                               A; Molecule type: DNA
A; Residues: 1-253 <SCH>
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C;Species: Colobus guereza
C;Date: 28-Oct-1996 #sequence_revision 07-Feb-1997 #text_change 13-Aug-1999
C;Accession: S53618; S71046
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A; Residues: 1-253 <SCW>
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A; Residues: 1-210, 'R', 212-253
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A; Residues: 1-253 <RES>
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160 QVYYRPVDQYSNQ 172
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A;Molecule type: DNA
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C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 major prion protein - hamadryas baboon
C;Species: Papio hamadryas (hamadryas baboon)
C;Date: 28-Oct-1996 #sequence_revision 07-Feb-1997 #text_change 13-Aug-1999
C;Accession: S53620; S71058
R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.
J. Mol. Biol. 245, 362-374, 1995
A;Reference number: S53614; MUID:95139066; PMID:7837269
A;Accession: S53620
A;Accession: S53620
A;Accession: S53620
Search completed: March 24, 2003, 17:24:14
Job time: 17.25 secs
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submitted to the EMBL Data Library, April 1994
A:Reference number: S71041
A:Accession: S71058
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A;Molecule type: DNA
A;Residues: 1-210, 'E', 212-253 <SCW>
A;Residues: 1-210, 'E', 212-253 <SCW>
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C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
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submitted to the EMBL Data Library, April 1994
A:Reference number: S71041
A:Accession: S71057
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                                                                                                    160 QYYYRPVDQYSNQ 172
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Pred. No. 0.0056;
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Maximum Match 100%
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58
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PRIO_MUSVI
PRIO_PIG
                                                    PRIO_CAMDR
PRIO_CAPHI
PRIO_FELCA
PRIO_SHEEP
PRP2_BOVIN
PRP2_TRAST
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PRIO_MACFA
                    PRIO_SAISC
PRIO_BOVIN
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                                                                     Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
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                                                   Prion;
                                                         PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                                   InterPro; IPR000817; Prion
16
216
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215
164
166
182
                                                   Glycoprotein;
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36	37	38	42	46	50	50	50	50	50	50	50
62.1	63.8	65.5	72.4	79.3	86.2	86.2	86.2	86.2	86.2	86.2	86.2
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POLG_HCVA	YE94_XYLFA	PRIO_TRIVU	PRIO_CANFA	PRIO_PANTR	PRIO_ODOHE	PRIO_CEREL	PRIO_SIGHI	PRIO_MESAU	PRIO_CRIMI	PRIO_CRIGR	PRIO_CALJA
P19712 hog cholera	Q9pd85 xylella fas				P47852 odocoileus			P04273 mesocricetu			P40247 callithrix

ALIGNMENTS

This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@iab-sib.ch). PRIO_CERAT STANDARD; PRT; 238 AA 095145; 095200; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 15-JUN-2002 (Rel. 41, Last annotation updat 15-JUN-2002 (Rel. 41, Last ann Ger Kuyl A.C., Dekker J.T., Goudsmit J.; Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases. -I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS. -I- SUBJULIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED Macaca sylvanus (Barbary ape). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae; Cercocebus aterrimus, and Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment). SEQUENCE FROM N.A. der Kuyl A.C., Dek Cercopithecinae; Cercocebus. NCBI_TaxID=36222, 9546; SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUTOLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONCIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC. update) ₽

15 238 238 215 199 166 182

BY SIMILARITY.

MAJOR PRION PROTEIN.

REMOVED IN MATURE FORM (BY SIMILARITY).

BY SIMILARITY.

BY SIMILARITY.

N-LINKED (GLCNAC. . .) (POTE

N-LINKED (GLCNAC. . .) (POTE

(BY SIMILARITY).

(POTENTIAL)

GPI-anchor; Repeat; Signal.

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                                                                                                                                                                                                                                                                                                                                                                                                                                                     der Kuyl A.C., Dekker J.T., Goudsmit J.;
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.

-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.

-I- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
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                                                                                                                                                                                            EMBL; U75383; AAB50630.1; -. HSSP; P04925; 1AG2. InterPro; IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Theropithecus gelada (Gelada baboon).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                             PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                       Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cercopithecinae; Theropithecus.
NCBI_TaxID-9565;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Major prion protein PRNP OR PRP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Q95270;
01-NOV-1997
                                        DOMAIN
                                                                CARBOHYD
                                                                            DISULFID
                                                                                                                                Prion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE FROM N.A.
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                                                                                                                                                                                                                                                                                                                                                  SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor.
DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND
ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOCICAL DISEASES KURUJ,
CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME
(GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE),
TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      QVYYRPVDQYSNQ 157
                                                                                                                                Brain; Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Similarity
9; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     (Rel.
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166
182
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61
69
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199
166
182
83

    Last sequence update)
    Last annotation update)
    precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).

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58
76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 51; DB Pred. No. 0.00 0; Mismatches
                                                                                                                                GPI-anchor; Repeat; Signal.
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                                     N-LINKED (GLCNAC. . .) (
N-LINKED (GLCNAC. . .) (
4 X 8 AA TANDEM REPEATS
                                                                           MAJOR PRION PROTEIN BY SIMILARITY.
                                                                                                      BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           5F59A3EBC3E3531B CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB 1;
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                                     'S OF P-H-G-G-G
                                        P-H-G-G-G-W-G-
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     RESULT 3
PRIO_AOTTR
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Best Local
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01-FEB-1995
01-NOV-1995
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REPEAT
NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;
"Prion protein gene variation among primates.";
J. Mol. Biol. 245:362-374(1995)
-i- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-i- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                   This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics in the European Bioinformatics in There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Major prion
    REPEAT
REPEAT
REPEAT
                                                                                                                                       SIGNAL
                                                                                                                                                                                                         InterPro; IPR000817; Prion. Pfam; PF00377; prion; 1. SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-95139066; PubMed-7837269; Schatzl H.M., Dacosta M., Taylor L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P40245;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PRIO_AOTTR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Aotus trivirgatus (Night monkey) (Douroucouli).
                                           REPEAT
                                                                    DOMAIN
                                                                                   CARBOHYD
                                                                                                                           CHAIN
                                                                                                                                                                    Prion;
                                                                                                                                                                               PROSITE;
                                                                                                                                                                                 PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                  EMBL; U08293; AAC50082.1; HSSP; P04925; 1AG2.
                                                                                                                                                                                                                                                                                         or send an email to license@isb-sib.ch).
                                                                                                 CARBOHYD
                                                                                                             DISULFID
                                                                                                                                                        NON_TER
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SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SUBCELLULAR LOCATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             "RODS"
                                                                                                                                                                   Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          238
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            (Rel. 31, Created)
(Rel. 31, Last sequence update)
(Rel. 32, Last annotation update)
protein precursor (PrP) (PrP27-30)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     69
    524
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                                                                                                                                                                   ycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         238
26104
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Chordata; Craniata; Vertebrata; Primates; Platyrrhini; Cebidae;
     59
67
75
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69.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Attached to the membrane by a GPI-anchor. IN HIGH QUANTITY IN THE BRAIN OF HUMANS A
BY SIMILARITY.

MAJOR PRION PROTEIN.

BY SIMILARITY.

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N-LINKED (GLCNAC. . .) (P

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Pred. No. 0.002;
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                                                                                                                                                                 Signal
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Euteleostomi;
                                                                       (POTENTIAL).
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MEDLINE-95139066; PubMed-7837269;
MEDLINE-95139066; PubMed-7837269;
Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;
*Prion protein gene variation among primates.";

*J. Mol. Biol. 245:362-374(1995).

*J. MOL. B
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01-FEB-1995 (Rel. 31, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Platyrrhini; Cebidae; Callicebinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRNP.
Callicebus moloch (Dusky titi).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P40248;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9523;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      152 QVYYRPVDQYSNQ 164
                                                                                                                                                      CARBOHYD
CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                     Pfam; PF00377; prion; SMART; SM00157; PRP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                          EMBL; U08312; AAC50100.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                                                                                                                                                                                           PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                         REPEAT
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                                                                                                                                                                                                                                                                                                 Prion;
                                                                                                                                                                                                                                                                                                                                                                                                                          InterPro; IPR000817; Prion.
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                                                  REPEAT
                                                                                                                           DOMAIN
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASES (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         "RODS"
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9; Conserv
                                                                                                                                                                                                                                                                                                      Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         76
239
239 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
     53
61
69
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STANDARD;
                                                                                                                                                                                                                                                                                                    Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                87.9%;
69.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               26246 MW;
  BY SIMILARITY.

MAJOR PRION PROTEIN.

BY SIMILARITY.

N-LINED (GLCNAC. ..) (POTENTIAL).

N-LINKED (GLCNAC. ..) (POTENTIAL).

5 X 8 AA TANDEM REPEATS OF P-H-G-G-W-G-Q.

1.
2.
3.
4.
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                                                                                                                                                                                                                                                                                                      GPI-anchor; Repeat; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2EFB77E354B7024A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        241 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 1; Length 239;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
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NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;
"Prion protein gene variation among primates.";
J. Mol. Biol. 245:362-374(1995).
-1- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-1- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED "RODS".
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Mandrillus sphinx (Mandrill) (Papio sphinx).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIO_MANSP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NCBI_TaxID=9561
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cercopithecinae; Mandrillus.
                                                                                                                                                                                                                                                                                                 entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                the European Bioinformatics Institute. There are no restrictions on use by non-profit institutions as long as its content is in no modified and this statement is not removed. Usage by and for commentations are not removed.
                                                                                                                                                                                                                                                                                                                                                                                          This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE-95139066; PubMed-7837269;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    153 QVYYRPVDQYSNQ 165
                                                                                                                                                                                                                                                                    EMBL; U08303; AAC50091.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
hes 9; Conserv
                                                                                                                                                                                                                           Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                            InterPro; IPR000817; Prion.
                                                                          CARBOHYD
                                                                                       DISULFID
                                                                                                         LIPID
                                                                                                                     PROPEP
                                                                                                                                      CHAIN
                                                                                                                                                    SIGNAL
                                                                                                                                                                   NON_TER
                                                                                                                                                                                Prion;
                                                                                                                                                                                                 PROSITE;
                REPEAT
                                           DOMAIN
                                                             CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 QVYYXPXDXYXNQ 13
                                                                                                                                                                                                                                                                                                                                                                                                                                        SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                PS00291; PRION_1; 1. PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             241 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STANDARD;
                                                                                                                                                                                Glycoprotein;
                                                                           223
>241
223
207
174
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69.2%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             26373 MW;

 Mismatches

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Pred. No. 0.002;
                                                                                                                                                                                    GPI-anchor; Repeat; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               C6D2013EE7CAEC93 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT;
                                             N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
5 X 8 AA TANDEM REPEATS OF P-H-G-G-G-W-G-
                                                                                          GPI-ANCHOR (BY SIMILARITY).
BY SIMILARITY.
                                                                                                                       REMOVED IN MATURE FORM (BY SIMILARITY).
                                                                                                                                         MAJOR PRION PROTEIN
                                                                                                                                                         BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              241 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1; Length 241;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Indels
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REPEAT

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Best Local
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                       DISULFID CARBOHYD
             CARBOHYD
                                                                                                                                    InterPro: IPRO00817; Prion.
Pfam; PF00377; Prion; 1
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
                                                               PROPEP
                                                                                         SIGNAL
                                                                                                            PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/order.
                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; U08292; AAC50081.1; HSSP; P04925; IAG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           P40250;
P40250;
O1-FEB-1995 (Rel. 31, Created)
O1-FEB-1995 (Rel. 31, Last sequence update)
16-CCT-2001 (Rel. 40, Last annotation update)
Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C).
                                                   TPID
                                                                                                                                                                                                                   EMBL; U08291; AAC50080.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cercopithecus aethiops (Green monkey) (Grivet), and Cercopithecus diana (Diana monkey). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cercopithecinae; Cercopithecus. NCBI_TaxID=9534, 36224;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CERAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REPEAT
NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
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REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RIO_CERAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              153 QVYYRPVDQYSNQ 165
                                                                                                                                                                                                                                                                                                                                                SUBCELIULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DECENERATIVE NEUTOLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MIKK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                               SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                1 QVYYXPXDXYXNQ 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     241 AA;
                                                                                               Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  241
26398 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           87.98;
69.28;
                                                                                          GPI-anchor; Repeat; Signal.
N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
                          REMOVED IN MATURE FORM (BY SIMILARITY).

GPI-ANCHOR (BY SIMILARITY).

BY SIMILARITY.
                                                                    MAJOR PRION PROTEIN.
                                                                                    BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 51; DB 1;
Pred. No. 0.002;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRT;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              E539D84E2E2B59DE CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           245 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1; Length 241;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            4; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Query Match
Best Local
                         DISULFID CARBOHYD
              CARBOHYL
                                                                                         CHAIN
                                                                                                      SIGNAL
                                                                                                                                                                          InterPro; IPR000817; Prion. Pfam; PF00377; Prion; 1. SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation. the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way endified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                          TIPID
                                                                       PROPEP
                                                                                                                                                                                                                       EMBL; U75386; AAB50625.1; -. EMBL; U75387; AAB50626.1; -. HSSP; P04925; 1AG2.
                                                                                                                                             PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                     Prion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     der Kuyl A.C., Dekker J.T., Goudsmit J.;
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
-1- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFEC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRIO_CERMO STANDARD; PRT; 246 AA. 095172; Q95173; 095173; 095173; 01-NOV-1997 (Rel. 35, Created) 01-NOV-1997 (Rel. 35, Last sequence update) 15-JUL-1998 (Rel. 36, Last annotation update) Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Cercopithecinae; Cercopithecus.
NCBI_TaxID=36226, 36227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cercopithecus mona, and Cercopithecus neglectus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      REPEAT
REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               152 QVYYRPVDQYSNQ 164
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DOMAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Local Similarity
les 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                           SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRE IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANUMALS INFECTED WITH THE DECEMERATIVE NEUTOLOGICAL DISEASES KURU, CREUTZFELLT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                   SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 QVYYXPXDXYXNQ 13
                                                                                                                                 Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        245 AA;
                                                                                                                              Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              51
         15
223
246
223
223
207
174
190
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     26885 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     59
67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        87.98;
                                                                                                                      GPI-anchor; Repeat; Signal.
                   MAJOR PRION PROTEIN.
REMOVED IN MATURE FORM (BY SIMILARITY).
GPI-ANCHOR (BY SIMILARITY).
BY SIMILARITY.
N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
                                                                                            BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Score 51; DB 1;
Pred. No. 0.0021;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ×
8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                D582B58E2726C99A CRC64;
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                                                                                                                                                     EMBL; U75388; AAB50627.1; -. HSSP; P04925; 1AG2.
InterPro; IPR000817; Prion.
Pfam; PF00377; Prion; 1.
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                           or send an email to license@isb-sib.ch).

EMBL; U75388; AAB50627 1.
                                                                                                                                                                                                                                                       This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss institute of Bioinformatics and the EMBL outstation the European Bioinformatics institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                             CARBOHYD
                                                       LIPID
                                                                                                                                PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                der Kuyl A.C., Dekker J.T., Goudsmit J.;
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.

-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
                  CARBOHYD
                                            DISULFID
                                                                    PROPEP
                                                                                                                      Prion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-NOV-1997 (Rel.
01-NOV-1997 (Rel.
01-NOV-1997 (Rel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1997 (Rel. 35, Last annotation update)
Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).
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REPEAT
REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NCBI_TaxID-27677;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cercopithecinae; Cercopithecus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Cercopithecus patas.
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Q95174;
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                                                                                                                                                                                                                                                                                                                                  SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUTOLOGICAL DISEASES KURG, CREUTZFELDT-JAKOB DISEASES (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                          SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS
                                                                                                                      Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Similarity
9; Conser
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69
                                                                                                                      Glycoprotein;
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35, Last sequence update)
35, Last annotation updat
    15
223
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69.2%;
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                                                                                                                  GPI-anchor; Repeat;
    N-LINKED
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                                     REMOVED IN MATURE FORM (BY SIMILARITY).

GPI-ANCHOR (BY SIMILARITY).

BY SIMILARITY.
              N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
                                                                            MAJOR PRION PROTEIN
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                                                                                             SIMILARITY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        8
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    TANDEM REPEATS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       DB 1;
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                                                                                                                Signal.
OF P-H-G-G-G-W-G-
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RESULT 9
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Best Local
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                                 DISULFID
CARBOHYD
CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A. Dekker J.T., Goudsmit J.;

der Kuyl A.C., Dekker J.T., Goudsmit J.;

submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.

-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.

HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
                                                                              LIPID
                                                                                                            CHAIN
                                                                                                                                                                                                                                                                                     use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entitles requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                      DOMAIN
                                                                                                                                                                   PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                 Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
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                                                                                              PROPEP
                                                                                                                          SIGNAL
                                                                                                                                       NON_TER
                                                                                                                                                        Prion;
                                                                                                                                                                                                                                             EMBL; U75385; AAB50628.1;
HSSP; P04925; lAG2.
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NCBI_TaxID=9531;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cercocebus torquatus atys (Red-crowned mangabey) (Sooty mangabey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Major prion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-NOV-1997
01-NOV-1997
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                                                                                                                                                                                                                               InterPro; IPR000817; Prion.
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                                                                                                                                                                                                                                                                                                                                                                                                                                  SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISSASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INTECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE),
                                                                                                                                                                                                                                                                                                                                                                                                         TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC. SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                       Brain;
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protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).
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61
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                                                                                                                                                     .ycoprotein;
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35, Last sequence update)
35, Last annotation updat
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26886
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69.2%;
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MAJOR PRION PROTEIN.

REMOVED IN MATURE FORM (BY SIMILARITY GPI-ANHOR (BY SIMILARITY).

BY SIMILARITY.

N-LINKED (GLCNAC. . .) (POTENTIAL).

N-LINKED (GLCNAC. . .) (POTENTIAL).

S X 8 AA TANDEM REPEATS OF P-H-G-G-G-G.
                                                                                                                                                  GPI-anchor; Repeat;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            0,:
                                                                                                                       BY SIMILARITY
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Pred. No. 0.
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0.0021;
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                                                                                        FORM (BY SIMILARITY).
                                                                                                                                                Signal
               OF P-H-G-G-G-W-G-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         0;
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REPEAT

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Best Local
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REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;
"Prion protein gene variation among primates.";
J. Mol. Biol. 245:562-374(1995).
-i- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-i- SUBBUNIT: PRP HAS A TENDENCY TO AGGREGATE XIELDING POLYMERS CALLED
                                              DISULFID
CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Platyrrhini; Cebidae; Cebinae; Cebus.
NCBI_TaxID=9515;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE
                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cebus apella (Brown-capped capuchin).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REPEAT
                                                                                                                                                                                                                                           EMBL; U08295; AAC50084.1; -. HSSP; P04156; 1E1G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIO_CEBAP
                                 CARBOHYD
                                                                                                                          SIGNAL
                                                                                                                                        Prion; Brain;
                                                                                                                                                     PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                   Pfam; PF00377; prion; 1
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                         or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE-95139066; PubMed-7837269;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P40249;
                                                                            CIPID
                                                                                                                                                                                                                             InterPro; IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      153 QVYYRPVDQYSNQ 165
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1 QVYYXPXDXYXNQ 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                          SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED MITH THE DEGENERATIVE HEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME
                                                                                                                                                                                                                                                                                                                                                                                                                  SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                             (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
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9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            246
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                 230
230
229
178
180
196
                                                                                                                                       Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STANDARD;
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76
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Score 51; DB Pred. No. 0.00 0; Mismatches
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MAJOR PRION PROTEIN.
REMOVED IN MATURE FORM (BY SIMILARITY).
GPI-ANCHOR (BY SIMILARITY).
BY SIMILARITY.
N-LINKED (GLCNAC. ..) (POTENTIAL).
N-LINKED (GLCNAC. ..) (POTENTIAL).
5 x 8 AA TANDEM REPEATS OF P-H-G-G-W-G-Q.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            F58679CBBEC5ADC7 CRC64;
                                                                                                                        SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1; Length 246;
0.0021;
                                                                                                                                       Repeat; Signal.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRIO_RABIT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT
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Best Local
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01-NOV-1997
01-NOV-1997
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REPEAT
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                                                    CARBOHYD
DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 species with apparent resistance to infection by prions.";
Gene 184:215-219(1997).
-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBBURIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oryctolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Major prion protein precursor (PrP) (PrP27-30) PRNP OR PRP.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                     InterPro; IPR000817; Prion. Pfam; PF00377; prion; 1. PRINTS; PR00341; PRION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -!- SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor.
-!- DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZEELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
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                                                                                                                                                                                                           SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                             entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                      CHAIN
                                                                                                                                                               SIGNAL
                                                                                                                                                                                     Prion; Brain;
                                                                                                                                                                                                                                                                                                                                                                     EMBL; U28334; AAC48697.1; -. HSSP; P10279; 1DWY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NCBI_TaxID=9986;
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                              DOMAIN
                                                                                                           CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -!- SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     "Characterization of a prion protein (PrP) gene from rabbit; a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Loftus B., Rogers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-97183665;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              159 QVYYRPVDQYSNQ 171
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9; Conser
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(Rel.
(Rel.
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178
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                                                                                                                                                                                     Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ξ
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       35, Created)35, Last sequence update)35, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PubMed=9031631;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            27579 MW;
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69.2%;
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BY SIMILARITY.
5 X 8 AA TANDE
Q.
                                                                                                                                                                                     GPI-anchor; Repeat; Signal
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                                                                               N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
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                                                                                                                                                               POTENTIAL
                              X 8 AA TANDEM REPEATS OF P-H-G-G-G-W-G-
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Mismatches
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RESULT 12
PRIO_COLGU
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Best Local Similarity
"hese 9; Conserv
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SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                        der Kuyl A.C., Dekker J.T., Goudsmit J.;
Submitted (CCT-1996) to the EMBL/GenBank/DDBJ databases.

-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.

-I- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
"RODS".
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               REPEAT
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REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Colobus guereza (Black-and-white colobus monkey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae; Colobinae;
                                                                                                                                            EMBL; U08297; AAC50086.1; -. EMBL; U75389; AAB50624.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                                                                  modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                             This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
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16-OCT-2001
                                                                                   PRINTS; PR00341; PRION. SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE OF 8-253 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., "Prion protein gene variation among primates.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    MEDLINE-95139066; PubMed-7837269;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Colobus.
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                              SIGNAL
                                                      PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                               InterPro;
                                                                                                               Pfam; PF00377; prion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     159 QVYYRPVDQYSNQ 171
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                                                                                                                                                                                                                                                                                                                                                              SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME
                                                                                                                                                                                                                                                                                                          (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC. SIMILARITY: BELONGS TO THE PRION FAMILY.
                                          Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Biol. 245:362-374(1995).
                                                                                                                              IPR000817; Prion.
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60
68
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(Rel. 31, Last sequence update)
(Rel. 40, Last annotation updat
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                                         Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ₹
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75
83
83
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27432 MW;
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precursor (PrP) (PrP27-30) (PrP33-35C).
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69.2%;
                                         GPI-anchor;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     5 <del>-</del> 4 2 -
            BY SIMILARITY.
MAJOR PRION PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 51; I
Pred. No. 0.
REMOVED IN MATURE FORM (BY SIMILARITY).
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                                       Repeat; Signal.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ₹
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01-FEB-1995 (Rel. 3
01-FEB-1995 (Rel. 3
15-JUL-1998 (Rel. 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CARBOHYD
CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gorilla gorilla gorilla (Lowland gorilla).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Gorilla.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        REPEAT
REPEAT
REPEAT
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DISULFID
                                                                                   This SWISS-PROT entry is copyright. It is produced through between the Swiss Institute of Bioinformatics and the Entrepean Bioinformatics Institute. There are no restructe by non-profit institutions as long as its content modified and this statement is not removed. Usage by and
                                                                                                                                                                                                                                                                                                              Cervenakova L., Brown P., Goldfarb L.G., Nagle J., Pettrone R., Rubenstein R., Dubnick M., Gibbs C.J., Gajdusek D.C.; "Infectious amylold precursor gene sequences in primates used for experimental transmission of human spongiform encephalopathy."; proc. Natl. Acad. Sci. U.S.A. 91:12159-12162(1994).

-I- FUNCTION: THE FUNCTION OF MERP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.; "Prion protein gene variation among primates."; J. Mol. Biol. 245:362-374(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE
EMBL; U08300; AAC50089.1; -. EMBL; U15166; AAA68633.1; -. HSSP; P04156; 1QLZ.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C).
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                                                        entities requires a license agreement (See http://www.isb-sib.ch/announce/
or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                  - I - SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. MEDLINE-95139066; PubMed-7837269;
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                                                                                                                                                                                          SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED MITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                               SIMILARITY: BELONGS TO THE PRION FAMILY.
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N-LINKED (GLCNAC. ..) (POTENTIAL).
5 x 8 AA TANDEM REPEATS OF P-H-G-G-W-G-Q.
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Pred. No. 0.0021;
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RA Deloukas P., Matthews L.H., Ashurst J., Burton J., Gilbert J.G.R.,
A Jones M., Stavrides G., Almeida J.P., Babbage A.K., Bagguley C.L.,
RA Balley J., Barlow K.F., Bates K.N., Beard L.M., Beare D.M.,
RA Beasley O.P., Bird C.P., Blakey S.E., Bridgeman A.M., Brown A.J.,
RA Buck D., Burrill W.D., Butler A.P., Carder C., Carter N.P.,
Chark S.Y., Clark S.Y., Clee C.M.,
Chapman J.C., Clamp M., Clark G., Clark L.N., Clark S.Y., Clee C.M.,
RA Chegy S., Cobley V.E., Collier R.E., Connor R.E., Corby N.R.,
Coulson A., Coville G.J., Deadman R., Dhami P.D., Dunn M.,
RA Clegg S., Cobley V.E., Collier R.E., Connor R.E., Corby N.R.,
RA Coulson A.G., Frankland J.A., Fraser A., French L., Garner P.,
RA Ellington A.G., Frankland J.A., Fraser A., French L., Garner P.,
RA Hammond S., Harley J.L., Heath P.D., Ho S., Holden J.L., Howden P.J.,
RA Huckle E., Hunt A.R., Hunt S.E., Jekosch K., Johnson C.M., Johnson D.,
RA Huckle E., Hunt A.R., Hunt S.E., Jekosch K., Johnson C.M., Lawlor S.,
RA Kay M.P., Kimberley A.M., King A., Kinghts A., Laird G.K., Lawlor S.,
RA Lehvaslatho M.H., Leversha M.A., Lloyd C., Lloyd D.M., Lovell J.D.,
RA Marsh V.L., Martin S.L., McConnachie L.J., McLay K., McMurray A.A.,
Milne S.A., Mistry D., Moore M.J.F., Mullikin J.C., Nickerson T.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIO_HUMAN
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01-NOV-1986
15-JUN-2002
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CONFLICT
SEQUENCE
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REPEAT
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
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Pfam; PF00377; prion; 1.
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GPI-ANCHOR (BY SIMILARITY)
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0.0021;
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Terwilliger J.D., Westaway D., Ott J.
"Linkage of a prion protein missense syndrome.";
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"Genetic and infectious prion dis-
"Genetic and 50:1129-1153(1993).
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Phillimore B.J.C.T., Prathalingam S.R., Plumb R.W., Ramsay H.,
Rice C.M., Ross M.T., Scott C.E., Sehra H.K., Shownkeen R., Sims S.,
Skuce C.D., Smith M.L., Soderlund C., Steward C.A., Sulston J.E.,
Swann R.M., Sycamore N., Taylor R., Tee L., Thomas D.W., Thorpe A.,
Tracey A., Tromans A.C., Vaudin M., Wall M., Wallis J.M.,
Whitehead S.L., Whittaker P., Willey D.L., Williams L., Williams S.A.,
Wilming L., Wray P.W., Hubbard T., Durbin R.M., Bentley D.R., Beck S.,
VARIANT FFI ASN-178.
MEDLINE=92195483; PubMed=1347910;
                                                                           Doh-Ura K., Tateishi J., Sasaki H., Kitamoto T., "Pro-->Leu change at position 102 of prion protei common but not the sole mutation related to Gerst
                                                                                                                       VARIANTS LEU-102; VAL-117 AND VAL-129.
WEDLINE-89392018; PubMed-2783132;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            "Amyloid protein of Gerstmann-Straussler-Scheinker disease (Indiana kindred) is an 11 kd fragment of prion protein with an N-terminal glycine at codon 58.";
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MEDLINE=94029646; PubMed=8105771;
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                                                                                                                                                                          Nature
                                                                                                                                                                                                                                        Hsiao K.,
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                                                                                                                                                                                                                                                                      VARIANT GSS LEU-102
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"Mutations and
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MEDLINE-20359708; PubMed-10900000;
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Liao Y.-C.J., Lebo R.V., Clawson G.A.,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 WEDLINE=20087216; PubMed=10618385;
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R., Wuethrich K.;
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233:364-367(1986).
                                               Biophys. Res.
                                                                                                                                                                                                                     Baker H.F., Crow T.J., Poer J.D., Westaway D., Ott
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                                             163:974-979(1989)
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Goldfarb L.G., Haltia M., Brown P., Nieto A., Kovanen J., McCombie W.R., Trapp S., Gajdusek D.C.;
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MEDLINE-93213314; PubMed-8461023;
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VARIANT GSS LEU-105.

MEDLINE-95213742; PubMed-7699395;

MEDLINE-95213742; PubMed-7699395;

MEDLINE-95213742; PubMed-7699395;

MICON Y., Yamada M., Hayakawa M., Shozawa T., Tanaka J., Matsushita M., Itoh Y., Yamada M., Hayakawa M., Shozawa T., Tanaka J., Matsushita M., Kitamoto T., Tatatshi J., Otomo E.;

Mitamoto T., Tatatshi J., Matsushita M., Kitamoto T., Tanaka J., Matsushita M., Mats
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                                                                                 MEDLINE-94142912; PubMed-7906019;
Inoue I., Kitamoto T., Doh-Ura K., Shii H., Goto I.,
"Japanese family with Creutzfeldt-Jakob disease with
mutation of the prion protein gene.";
Neurology 44:299-301(1994).
                                                                                                                                                                                                                                                                                           clinicopathological study.";
J. Neurol. Sci. 127:77-86(1994).
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Tateishi J., Otomo E., Hayakawa M., Tanaka J., Matsushita M.,
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MEDLINE-94077414; PubMed-7902972;
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   VARIANT CJD LYS-200.
MEDLINE-94316708; PubMed-7913755;
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MEDLINE-95139066; PubMed-7837269;

Schatzl H.M., Dacosta M., Taylor L., Cohen F.E.

"Prion protein gene variation among primates.";
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1- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
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Mammalia; Eutheria; Primates; Catarrhini;
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This SWISS-PROT entry is copyright. It is produced through a collaboration
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                              (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC. SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                              SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CDD), GERSTMANN-STRAUSSLER SYNDROME
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212	204	204	204	185	141	285	264	257	256	256	256	256	256	256	256	256	256	256	254	254	254	253	253	253	253	245	235	233
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ALIGNMENTS

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40.	 Ma Ma	SO N N N N N N N N N N N N N N N N N N N
1 QVYYXPXXXXXNQ 13 1 93 QVYYKPVDQYSNQ 105	Query Match 87.9%; Score 51; DB 6; Length 124; Best Local Similarity 69.2%; Pred. No. 0.006; Matches 9; Conservative 0; Mismatches 4; Indels 0; Gaps	Q9TU20 Q9TU20; Q9TU20; Q9TU20; Q9TU20; Q1-MAY-2000 (TrEMBLrel. 13, Created) Q1-MAY-2000 (TrEMBLrel. 13, Last sequence update) Q1-MAY-2000 (TrEMBLrel. 19, Last sequence update) Prion Protein (Fragment). PRE: Varecia variegata variegata. Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalla; Eutheria; Primates; Strepsirhini; Lemuridae; Varecia. MCBI_TaxID-87289; [1] SEQUENCE FROM N.A. Gilch S., Schatzl H.M.; UDUSUAL PRION Cotarepeat structure of the highly BSE- susceptible lemur monkey."; Submitted (AUG-1999) to the EMBL/GenBank/DDBJ databases. EMBL; AF177293; AAD54335.1; HSSP; P04925; 1AG2. InterPro; IPR000817; Prion. PFiam; PF00377; prion; 1. PRINTS; PR00141; PRION. SMART; SM00157; PRP; 1. PRON_TER 124 124 SEQUENCE 124 AA; 13436 MW; CC2C8A5A855A7C94 CRC64;

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RESULT 3
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                                  Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Schwarz T.F., Werner T., Schwatzl H.M.;
"Analysis of 27 mammalian and 9 avian PrPs reveals high of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
EMBL; AF117309; AAD19980.1; -.
HSSP; P10279; IDWY.
InterPro; IPR002395; Kininogen.
InterPro; IPR000817; Prion.
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InterPro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
PRINTS; PR0034; KININOGEN.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
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01-MAY-1999 (TrEMBLrel. 10,
01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                        MEDLINE-99303687; PubMed-10373359;
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181 AA;
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69.2%;
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Pred. No. 0.
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Ruminantia; Pecora; Bovoidea
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097912;
01-MAY-1999 (TrEMBLrel. 1
01-MAY-1999 (TrEMBLrel. 1
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        Bison bonasus (European bison).
Eukaryota; Metazoa; Chordata; Crania
Mammalia; Eutheria; Cetartiodactyla;
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PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Canis lupus (Gray wolf).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
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01-MAY-1999 (TrEMBLrel. 10,
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PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1
.PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
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Pred. No. 0.
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Pred. No. 0.
                   Craniata; Vertebrata; Euteleostomi;
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ion protein.";
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Query Match
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PRINTS; PR00334; KININGEN.

PRINTS; PR00341; PRION.

SMART; SM00086; PAC; 1.

SMART; SM00157; PRP; 1.

PROSITE; PS00291; PRION_1; 1.

PROSITE; PS00706; PRION_2; 1.
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01-MAY-1999 (TrEMBLrel. 10,
01-MAY-1999 (TrEMBLrel. 10,
01-JUN-2002 (TrEMBLrel. 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDILINE-99303687; PubMed-10373359; Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schatzl H.M.; Schwarz T.F., Werner T., Schatzl H.M.; "Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein."; J. Mol. Biol. 289:1163-1178(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                      097908
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                          Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schatzl H.M.; "Analysis of 27 mammalian and 9 avian PrPs reveals high conservation
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                HSSP; P10279; 1DWY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           EMBL; AF117328; AAD19999.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9902;
                                                                                                                                                                                                                                                                                                                                                  Capra nubiana (Nubian ibex).
                                                                                                                                                                                                                                                                                                                                                                         Prion protein (Fragment).
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                       SEQUENCE
                                                           PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                                                Pfam; PF00377; prion; 1. PRINTS; PR00334; KININOGEN. PRINTS; PR00341; PRION.
                                                                                                                                                InterPro;
                                                                                                                                                                                      of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
                                                                                                                                                                                                                                              MEDLINE-99303687; PubMed-10373359;
                                                                                                                                                                                                                                                                         SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                               Bovidae; Caprinae; Capra.
NCBI_TaxID-72543;
                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  128
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                                                                                                                                                                         AF117319; AAD19990.1;
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9; Conservative
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IPR000817; Prion.
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21674 MW;
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  87.98;
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                        DB0634A43B4DB77F CRC64;
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Matches 9
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Matches 9; Conserv
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**Rhalysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein.**;

J. Mol. Biol. 289:1163-1178(1999).

EMBL; AF113943; AAD13291.1; -.

HSSP; P10279; IDWY.
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Eukaryota; Metazoa;
Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-MAY-1999 (TrEMBLrel. 01-JUN-2002 (TrEMBLrel.
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MEDLINE=99303687; PubMed=10373359;
Wopfner F., Weidenhofer G., Schnei
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prion protein (Fragment).
                                                                    SEQUENCE FROM N.A.
MEDLINE-99303687; PubMed-10373359;
MEDLINE-99303687; PubMed-10373359;
Wopfner F., Weidenhofer G., Schneider R., von Brunn A.,
Schwarz T.F., Werner T., Schatzl H.M.;
"Analysis of 27 mammalian and 9 avian PrPs reveals high
of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
                                                                                                                                                                                                                                                                                                                                                                                                    Camelus dromedarius (Dromedary) (Arabian camel).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Tylopoda; Camelidae; Camelus
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    01-MAY-2000 (TIEMBLIEL 13, 01-MAY-2000 (TIEMBLIEL 13, 01-JUN-2002 (TIEMBLIEL 21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Camelus dromedarius
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PRINTS; PR00341; PRION.
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                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=9838;
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       P10279; 1DWY
                                         AF113940; AAD13288.1; -.
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202 AA;
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21860 MW;
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69.28;
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                                                                                                                                                                                                                             Gilch S.,
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                                                                                                                                                                                                                                                                                                          InterPro; IPR002395; Kininogen.
InterPro; IPR000817; Prion.
Pfam; PF00377; Prion; 1.
PRINTS; PR00334; KININOGEN.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        InterPro; IPR000817; Prion.
Pfam; PF00377; Prion; 1.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
Canis familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                            01-MAY-2000 (TrEMBLrel. 13, 01-MAY-2000 (TrEMBLrel. 13, 01-DEC-2001 (TrEMBLrel. 19, Prion protein (Fragment).
                                                                                 01-MAY-2000
                                                                                              Q9TV04;
                                                                                                       Q9TV04
                                                                                                                                                                                                                                                                                                                                                                               O'Rourke K.I., Miller M.W., Wild M.A., Williams E.S.;
"PrP gene of pronghorn antelope (Antilocapra americana) contains octapeptide repeats.";
submitted (SEP-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF090852; AAC43030.1; -...
HSSP; P10279; IDWY.
                                                                                                                                                                                                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Antilocapra americana (Pronghorn).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Antilocapridae; Antilocapra.
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NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                    PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                   TISSUE-BRAIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID-9891;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  01-NOV-1998 (TrEMBLrel. 08, Created)
01-NOV-1998 (TrEMBLrel. 08, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
Prion protein (Fragment).
                                                                                                                                                               148
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   126 QVYYKPVDQYSNQ 138
                                                                                                                                                                          1 QVYYXPXDXYXNQ 13
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                                                                                                                                                               QVYYRPVDQYSNQ 160
                                                                                                                                                                                                          Similarity
9; Conserv
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                                                                                                                                                                                                                                                     211
211 AA;
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209 AA;
                                                                                                                                                                                                          Conservative
                                                                                                     PRELIMINARY;
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69.2%;
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69.28;
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                                                      Last annotation update)
                                                                  Last sequence update)
                                                                                                                                                                                                                 Score 51; DB 6; Length 211; Pred. No. 0.01;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0
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Pred. No.
                                                                                                     PRT;
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                                                                                                                                                                                                                                                    B8E147AADF9A6752 CRC64;
                                                                                                                                                                                                        Mismatches
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                                                                                                     213 AA.
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                                                                         RESULT 11
Q9TV03
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Matches 9
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                                                                                                                                        EMBL; AFI13938; AAD12062.1; -
HSSP; P04925; 1AG2.
InterPro; IPRO00817; Prion.
Pfam; P800377; Prion; 1.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
                                                                                 VARIANT
NON_TER
                                                                                                          PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                  Q9TV03;
                                                                                                                                                                                                              MEDLINE-99303687; PubMed-10373359; Wopfner F., Weldenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schatzl H.M.; R., Schwarz T.F., Werner T., Schatzl H.M.; Panalysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein."; J. Mol. Biol. 289:1163-1178(1999).
                                                                                                                 PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                                                                                                                                                                                                                                                                           Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Canivora; Fissipedia; Canidae;
                                                                        SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                            01-MAY-2000 (TrEMBLrel.
01-MAY-2000 (TrEMBLrel.
01-DEC-2001 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NON_TER
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PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                     STRAIN-BREED DACHSHUND;
                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                   Prion protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           of flexible regions of the prion protein."
J. Mol. Biol. 289:1163-1178(1999).
EMBL; AFIL3937, AAD12061.1;
HSSP: P04925; 1AG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schatzl H.M., Schwarz Schwarz Schwarz Schwarz H.M., Weiden Pres reveals high conservation "Analysis of 27 mammalian and 9 avian Pres reveals high conservation"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         123 QVYYRPVDQYSNQ 135
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           SEQUENCE FROM N.A. MEDLINE-99303687; PubMed-10373359;
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                                    Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Local Similarity 69.7 tes 9; Conservative
1 QVYYXPXDXYXNQ 13
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                         Similarity
9; Conserv
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213 AA;
                       Conservative
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22997 MW;
                                                                      23167 MW;
                                 87.9%;
69.2%;
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69.28;
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                                Score 51; DB 6; Pred. No. 0.011;
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Pred. No. 0.01;
                     Mismatches
                                                                    551B7669ABD4C6DF CRC64;
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                                             Length 214;
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HMSSP; P10279; LDWY:
InterPro; IPR002395; Kininogen.
InterPro; IPR001610; PAC.
InterPro; IPR009817; Prion.
Pfem; PF00377; Prion; 1.
PRINTS; PR00344; KININOGEN.
PRINTS; PR00344; FRION.
SMART; SM00086; PAC; 1.
SMART; SM00157; PRP; 1.
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01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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"Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
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                                                                                                                                                                                                                                                                                                                                                            Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoldea;
Bovidae; Bovinae; Bos.
NCBI_TaxID-9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAY-2000 (TrEMBLrel. 13, 01-MAY-2000 (TrEMBLrel. 13, 01-JUN-2002 (TrEMBLrel. 21,
MEDIJINE-9303687; PubMed-10373359; Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schatzl H.M.; Schwarz T.F., Werner T., Schatzl H.M.; Analysis of 27 mammalian and 9 avian Prps reveals high conservation of flexible regions of the prion protein."; J. Mol. Biol. 289:1163-1178(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Q9TV00;
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                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     136 QVYYRPVDQYSNQ 148
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity es 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  1 QVYYXPXDXYXNQ 13
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215 AA;
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23182 MW;
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69.2%;
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Pred. No. 0.011;
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097907;

PRELIMINARY;

226 AA.

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RESULT 14
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RESULT 15
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Best Local Similarity
Matches 9; Conserv
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097910;
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InterPro; IPRO00817; Prion.
Pfam; PF00377; prion; 1.
PRINTS; PR00341; PRION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               EMBL; AF117327; AAD19998.1; HSSP; P10279; 1DWY.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SMART; SM00086; PAC; 1.
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hippotragus niger (Sable antelope).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                         Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Glich S., Schwarz T.F., Werner T., Schatzl H.M.; Schwarz T.F., Werner T., Schatzl H.M.; "Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein."; J. Mol. Biol. 289:1163-1178(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prion protein (Fragment).
                                                                                                                                                                                                                               Interpro; IPR002395; Klninogen.
Interpro; IPR000817; Prion.
Pfam; PF00377; prion; 1
PRINTS; PR00334; KININOGEN.
PRINTS; PR00341; PRION.
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EMBL; AF117323; AAD19994.1; -.
                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-99303687; PubMed-10373359;
                                                                                                                                                                                                                                                                                                                                                                                                                                       Bovidae; Hippotraginae; Hippotragus.
NCBI_TaxID-37189;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 131 QVYYRPVDQYSNQ 143
                                                                                                                                                                                           PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                                                                                                                                                                       SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                HSSP; P10279; 1DWY.
                                                                                                                                                                                                                                                                                                                                                                                                                   SEQUENCE FROM N.A.
                                                                                                                                                     SEQUENCE
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                                                   136 QVYYRPVDQYSNQ 148
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                                                                          1 QVYYXPXDXYXNQ 13
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223 AA;
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                                                                                                    Conservative
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23425 MW;
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24172 MW;
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                                                                                                                 87.9%;
69.2%;
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                                                                                                                   Score 51; DB 6; pred. No. 0.011;
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                                                                                                                                                         77A95AC13080F416 CRC64;
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Search completed: March 24, 2003, 17:22:11 Job time: 28.4375 secs
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                                                                                                                          Ouery Match 87.9%; Score 51; DB 6; Length 226; Best Local Similarity 69.2%; Pred. No. 0.011; Matches 9; Conservative 0; Mismatches 4; Indels
                                                                                                                                                                                       InterPro; IPRO02395; Kininogen.
InterPro; IPRO02395; Kininogen.
InterPro; IPRO0817; prion.
Pfam; PP00377; prion; 1.
PRINTS; PR00334; KININOGEN.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00291; PRION_2; 1.
NON_TER 1 1
VARIANT 209 209 R ->
NON_TER 226 AA; 24384 MW; D8
SEQUENCE 226 AA; 24384 MW; D8
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Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S.,
Schwarz T.F., Werner T., Schatzl H.M.;
"Analysis of 27 mammalian and 9 avian PrPs reveals high conservation
of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
EMBL; AF117313; AAD19984.1; -.
HSSP; P10279; IDMY.
HSSP; P10279; IDMY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gazella subgutturosa (Goltred gazelle).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovoidae; Antilopinae; Gazella.
                                                                    141 OVYYRPVDOYSNO 153
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last seq.
01-JUN-2002 (TrEMBLrel. 21, Last anno
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TISSUE-PBL;
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                                                                                       1 QVYYXPXDXYXNQ 13
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226
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Result
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Maximum Match 100%
Listing first 45 summaries
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score greater than or equal to the score of the result being printed,
and is derived by analysis of the total score distribution.
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seq length: 2000000000
      protein search, using sw model
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Match
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Gapop 10.0 , Gapext 0.5
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58
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    /cgn2_6/ptodata/2/iaa/5A_COMB.pep:*
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/cgn2_6/ptodata/2/iaa/PCTUS_COMB.pep:*
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US-09-076-721-45
US-08-244-701B-16
US-08-244-701B-18
US-09-076-721-17
US-09-076-721-18
US-09-076-721-18
US-09-076-721-18
US-09-076-721-13
US-09-076-721-13
US-09-076-721-14
US-09-076-721-14
US-09-076-721-14
US-09-076-721-14
US-09-076-721-12
US-08-556-823-10
US-08-556-823-10
US-08-60-626-8
US-09-031-168-8
US-09-031-158-2
US-09-031-158-2
US-09-031-158-8
US-09-031-158-8
US-09-031-158-8
US-09-031-158-8
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US-08-244-701B-45
US-09-076-721-44
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0 7	Query Best Match	SEGULT S-08-2 Seque Seque Pateu Pateu App App TI I O CO CO INFO SEE SEE SEE SEE SEE SEE SEE SEE SEE SE		. 4444433333333333333333333333333333333
7 1 QVYYXPXDXYXNQ 13 	9%; Score 51; DB 1; Len 2%; Pred. No. 0.0016; 0; Mismatches 4;	- O8 - NE T A C C C C C N T T A A C C C C N T T A A C C C C C C C C C C C C C C C C	ALIGNMENTS	28 51 87.9 253 4 US-09-550-374-2 29 51 87.9 254 1 US-08-242-188-1 30 51 87.9 254 1 US-08-60-626-7 31 51 87.9 254 1 US-08-692-892-1 33 51 87.9 254 1 US-08-692-892-1 34 51 87.9 254 2 US-08-868-162A-21 35 51 87.9 254 4 US-09-128-450-19 36 51 87.9 254 4 US-09-128-450-19 37 51 87.9 254 4 US-09-128-450-28 38 51 87.9 254 4 US-09-128-450-28 38 51 87.9 254 4 US-09-128-494-19 40 51 87.9 254 4 US-09-823-494-19 40 51 87.9 254 4 US-09-823-494-19 41 51 87.9 255 1 US-08-262-10 42 51 87.9 255 1 US-08-262-10 43 51 87.9 255 1 US-08-60-626-10 45 87.9 255 1 US-08-60-626-10
	gth 27; Indels 0;			Sequence Sequence
	Gaps			2, App 1, App 1, App 1, App 1, App 1, App 21, App 21, App 11, App 11, App 11, App 11, App 12, App 14, App 14, App 14, App 10, App
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US-08-244-701B-45
                                                                                                                                                                                                               Sequence 44, Application US/09076721 Patent No. 6379905 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                       Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELEFAX: (212) 869-8864/97.
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 45:
SEQUENCE CHARACTERISTICS:
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                         CORRESPONDENCE ADDRESS:
ADDRESSEE: Pennie & Edmonds
                                                                                                                                  APPLICANT: Mee, Roger P.
TITLE OF INVENTION: Fragments of Prion Proteins
NUMBER OF SEQUENCES: 67
                                                                                                                                                                     APPLICANT: F1shl
APPLICANT: Robso
APPLICANT: Mee,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY: linear MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (212) 790-9090
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          LENGTH: 27 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                   ZIP:
                                                                                            STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NAME: Fanucci, Allan A. REGISTRATION NUMBER: 30, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE: 02-JUN-1994
CLASSIFICATION: 436
                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity es 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/244,701B FILING DATE: 02-JUN-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STREET:
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                                                                            New York
                                                               New York
                                                                                          1155 Avenue of the Americas
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                                              U.S.A.
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                                                                                                                                                                                    Robson, Barry
                                                                                                                                                                                                  Fishleigh, Robert V.
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SYSTEM: PC-DOS/MS-DOS
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69.2%;
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                      0.0016;
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Query Match
Best Local Similarity
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US-09-076-721-45
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                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/076,721
                                                             FILING DATE:
ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 8080-007
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                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICANT: Mee, Roger P.
TITLE OF INVENTION: Fragments of Prion Proteins
NUMBER OF SEQUENCES: 67
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LENGTH: 27 amino acids
                              TELEPHONE: (212) 790-9090
                                                                                                                                              PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT:
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                                                                                                                                                                                FILING DATE:
CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                           STREET: 1155 A. CITY: New York
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TELEFAX: 66141 PENNIE
TELEX: 66141 PENNIE
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                                                                                                                                                                                                                                                                                                                                                          STATE: New York
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         FILING DATE:
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66141 PENNIE
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1155 Avenue of the
                  (212) 869-8864/9741
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SYSTEM: PC-DOS/MS-DOS
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linear
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                                                                                                                                                US 08/244,701
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                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 16:
SEQUENCE CHARACTERISTICS:
LENGTH: 28 amino acids
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Best Local Similarity 69.2%;
Matches 9; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0,
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                   ATTORNEY/AGENT INFORMATION: NAME: Fanucci, Allan A.
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MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                          MOLECULE TYPE:
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                                                     NAME/KEY:
   OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                       OTHER INFORMATION: OTHER INFORMATION:
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STRANDEDNESS: sir
TOPOLOGY: linear
                                                                                                                                       OTHER INFORMATION:
                                                                                                                                                                        NAME/KEY:
                                                                                                                                                                                                                              TOPOLOGY: 11
                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Fanucci, Allan A. REGISTRATION NUMBER: 30,256
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STATE: New York
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           STREET:
                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 8080-007
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1155 Avenue of the Americas
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Mee, Roger P.
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linear
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                                                                                                 /label= X
/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
/label= Y
/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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Pred. No. 0.0016;
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                                                         Best Loc
Matches
                                                                                     Query Match
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                                                                                                                                                                                                                                                                                                                                                                                              TELEFAX: (212) 869-886
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                              FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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                                                                                                                               LOCATION: 28
OTHER INFORMATION:
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide
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nes 9; Conserv
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OTHER INFORMATION:
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CITY: New York
STATE: New Yor
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                                                                                                                                                                                                                                                         OTHER INFORMATION:
                                                                                                                                                                                                                                                                         LOCATION:
                                                                                                                                                                                                                                                                                                                                                                TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                FILING DATE: 02-JUN-1994
CLASSIFICATION: 436
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6 QVYYRPVDRYSNQ 18
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(212) 869-8864/9741
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                                                         Conservative
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SYSTEM: PC-DOS/MS-DOS
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                                                                       87.9%;
69.2%;
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                                                                                                                               /label= y
/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s) "
                                                                                                                                                                                                                    /label= x
/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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                                                     Score 51; DB 1;
Pred. No. 0.0017;
0; Mismatches
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Pred. No. 0.0017;
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US-09-076-721-16
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                                   Sequence 16, Application US/09076721
Patent NO. 6379905
GENERAL INFORMATION:
APPLICANT: Fishleigh, Robert V.
APPLICANT: Robson, Barry
APPLICANT: Mee, Roger P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 18, Application US/08244701B Patent No. 5773572
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Best Local Similarity
Matches 9; Conserv
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TELEX: 66141 PENNIE
INFORMATION FOR SEI IN NO:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPHONE: (212) 869-8864/9741
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NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
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       FITLE OF INVENTION:
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OTHER INFORMATION:
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SOFTWARE: PatentIn Release #1.0, Version #1.25
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                               Robson, Barry
Mee, Roger P.
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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Fragments of Prion Proteins
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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Pred. No. 0.0017;
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                                                                                                                                                                                                                                                                                                                    Sequence 17, Application US/09076721
Patent No. 6379905
GENERAL IMPORMATION:
APPLICANT: Fishleigh, Robert V.
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ACTERISTICS:

ACTER
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APPLICANT: Mee, Roger P.
TITLE OF INVENTION: Fragments of Prion Proteins
NUMBER OF SEQUENCES: 67
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MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0
                                                                                                                                     CORRESPONDENCE ADDRESS:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
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NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
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OTHER INFORMATION:
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APPLICATION NUMBER:
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CITY: New York
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STREET: 1155 Avenue of the Americas
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                                        New York
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                                                           E: Pennie & Edmonds
1155 Avenue of the Americas
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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Pred. No. 0.0017;
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US-09-076-721-17
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Patent No. 6379905
GENERAL INFORMATION:
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Best Local Similarity
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OPERATING SYSTEM: PC-DOS/MS-DOS
SOPTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                              APPLICANT: Robson, Barry
APPLICANT: Wee, Roger P.
TITLE OF INVENTION: Fragme
NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPHON: (212) 869-8864/9741
TELEPAX: (6141 PENNIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 801
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION DATA:
APPLICATION NUMBER: U
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LENGTH: 28 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     FEATURE:
COMPUTER READABLE FORM:
MEDIUM TYPE: F10PPY d1sk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                           APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FILING DATE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 OTHER INFORMATION: /label= Y OTHER INFORMATION: /note= "Y may be absent or present independently OTHER INFORMATION: of X and denotes one or more amino acid(s) "
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          OTHER INFORMATION: /label= X OTHER INFORMATION: /note= "X may be absent or present independently OTHER INFORMATION: of Y and denotes one or more amino acid(s)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NAME/KEY: Modified-site LOCATION: 28
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                                                                       COUNTRY: U
ZIP: 10036
                                                                                                           STREET: 1155 A
CITY: New York
STATE: New Yor
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                                                                                                                                                                 ADDRESSEE:
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10036
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                                                                                                             New York
                                                                                                                                              E: Pennie & Edmonds
1155 Avenue of the Americas
                                                                                                                                                                                                                                                                       Fishleigh, Robert V.
                                                                                          U.S.A.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             87.9%; Score 51; DB 4; 69.2%; Pred. No. 0.0017;
                                                                                                                                                                                                     Fragments of Prion Proteins 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US 08/244,701
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                                                                                                                                                                                                                                                                                                                                                                                           Sequence 13, Application US/08244701B Patent No. 5773572 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-08-244-701B-13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEX: 66141 PENNIE INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/09/076,721
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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           COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMBER: US/08/244,701B
FILING DATE: 02-JUN-1994
CLASSIFICATION: 436
CHASSIFICATION: 436
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OTHER INFORMATION: /
OTHER INFORMATION: /
OTHER INFORMATION: C
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                                                                                                                                                                                                                                                                                         CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                                                                                                 APPLICANT:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: Fanucci, Allan A. REGISTRATION NUMBER: 30,256
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: /label= Y OTHER INFORMATION: /note= "Y may be absent or present independently OTHER INFORMATION: of X and denotes one or more amino acid(s)"
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
STRANDEDNESS: sir
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                                                                                                                                                                  ZIP: 10036
                                                                                                                                                                                                                    CITY: New York
STATE: New York
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linear
                                                                                                                                                                                                                                                        1155 Avenue of the Americas
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69.2%;
                                                                                                                                                                                                                                                                                                              Fragments of Prion Proteins 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /label= X
/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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US-08-244-701B-14
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OTHER INFORMATION:
OTHER INFORMATION:
US-08-244-701B-13
                                            COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/244,701B
FILING DATE: 02-UN-1994
CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 8080-007
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEPHONE: (212) 869-8864/9741
TELEPHONE: (212) 869-8864/9741
TELEFAX: (212) 869-8864/97.
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO: 14:
SEQUENCE CHARACTERISTICS:
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Patent No. 5773572
GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 13:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
                                                                                                                                                                                                                                                                                                                                                                                                                                                          NUMBER OF SEQUENCES: 6
CORRESPONDENCE ADDRESS:
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APPLICANT: Mee, Roger P.
TITLE OF INVENTION: Fragi
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REFERENCE/DOCKET NUMBER: 8080
TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
                                                                                                                                                                                                                                                                                                                                                                               STREET: 1155 Ave
CITY: New York
STATE: New York
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nes 9; Conserv
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OTHER INFORMATION:
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         87.9%; Score 51; DB 1; 69.2%; Pred. No. 0.0019;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Fragments of Prion Proteins 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      /label= Y
/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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/note= "x may be absent or present independently
of Y and denotes one or more amino acid(s)"
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RESULT 13
US-08-244-701B-15
ADDITIONAL TO THE PROPERTY OF THE 
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                                                                                                                                                                                                                                                                                                 TELEFAX: (212) 869-886
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICATION NUMBER: US/08/:
APPLICATION NUMBER: US/08/:
FILING DATE: 02-UUN-1994
CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan A.
NAME: Fanucci, Allan A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Matches
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                                                                                                                        MOLECULE TYPE: peptide FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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APPLICANT:
APPLICANT:
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STRANDEDESS: "
TOPOLOGY:
OLECUT
OTHER INFORMATION: /label- x
                                                                                                                                                                                                      LENGTH: 31 amino acids
TYPE: amino acid
STRANDEDNESS: single
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                                                                          NAME/KEY: Modified-site
                                                                                                                                                                                TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REGISTRATION NUMBER: 30, REFERENCE/DOCKET NUMBER:
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OTHER INFORMATION: /label= Y
OTHER INFORMATION: /note= "Y may be absent or present independently
OTHER INFORMATION: of X and denotes one or more amino acid(s)*
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            87.9%;
69.2%;
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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Pred. No. 0.0019;
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Query Match
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INFORMATION FOR SEQ ID NO:
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tent No. 6379905
                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 31 amino acids
                                                                                                                                                                                                                                                                                                                         NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 80E
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/09/076,721
                                                                                                                                                                                                                                                                                                                                                                                                                                            CLASSIFICATION:
PRIOR APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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                                                   FEATURE:
                                                                                                                                                                OLECULE TYPE: peptide
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                                                                                                                               NAME/KEY: Modified-site
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                                                                                                                                                                                               LENGTH: 31 amino acids
TYPE: amino acid
STRANDEDNESS: single
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OPERATING SYSTEM: PC-DOS/MS-DOS
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OTHER INFORMATION:
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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    of Y and denotes one or more amino acid(s)"
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Pred. No.
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US-09-076-721-14
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US-09-076-721-14
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 Query Match
Best Local Similarity
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GENERAL INFORMATION:
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Best Local
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APPLICANT:
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NAME: FANUCCI, Allan A.
REGISTRATION UNMBER: 30,256
REFERENCE/DOCKET NUMBER: 801
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MEDIUM TYPE: Floppy disk
COMPOTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
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PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                        FEATURE:
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LENGTH: 31 amino acids
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                                                                  OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                    LENGTH: 31 amino acids
TYPE: amino acid
STRANDEDNESS: single
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STATE: New York
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Local Similarity 69.2%;
es 9; Conservation
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1155 Avenue of the Americas
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 87.9%;
69.2%;
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                                                           /label= Y
/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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Score 51; DB 4;
Pred. No. 0.0019;
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Pred. No. 0.0019;
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              Length 31;
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Search completed: March 24, 2003, 17:23:05 Job time: 13.4583 secs

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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                Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Score
                protein search, using sw model
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query
Match
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58
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              Published_Applications_AA: *
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/cgn2_6/ptodata/1/pubpaa/US06_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US06_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US07_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US07_NEW_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US07_PUBCOMB.pep:*
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/cgn2_6/ptodata/1/pubpaa/US09_PUBCOMB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US10_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/US60_NEW_PUB.pep:*
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                   US-09-745-003-7
US-09-745-003-7
US-09-745-003-10
US-09-745-003-12
US-09-745-003-12
US-09-745-003-12
US-09-943-906-2
US-09-943-906-2
US-10-106-574-6
US-10-106-574-7
US-10-106-
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Sequence 3, Appli
Sequence 57, Appli
Sequence 2, Appli
Sequence 5, Appli
Sequence 6, Appli
Sequence 7, Appli
Sequence 1, Appli
Sequence 1, Appli
Sequence 4, Appli
Sequence 2, Appli
Sequence 3, Appli
Sequence 3, Appli
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Sequence 3, Appli
Sequence 2, Appli
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Sequence 4, Appli
Sequence 7, Appli
Sequence 9, Appli
Sequence 10, Appl
Sequence 12, Appl
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sequence souro, A		,	7		ń		Sequence 7, Appli			554	Sequence 6, Appli		12469,	3	5684	346	Sequence 4, Appli	Sequence 4, Appli	Sequence 244, App	Sequence 250, App	440	0.	œ :		e 11	Sequence 2, Appli

ALIGNMENTS

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RESULT 2
US-09-147-761-4
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US-09-939-780-4
Sequence 4, Application US/09147761
Patent No. US20010010918A1
GENERAL INFORMATION:
APPLICANT:
TITLE OF INVENTION: IMMUNOLOGIC
TITLE OF INVENTION: ENCEPHALOPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 4, Application US/09939780
Patent NO. US20020168689A1
GENERAL INFORMATION:
APPLICANT: O'Connor, Michael
TITLE OF INVENTION: Immunological Assay for Spongiform Encephalopathies
FILE REFERENCE: 50002US
                                                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PatentIn Ver. SEQ ID NO 4
LENGTH: 46
                                                                                                                                                                                                                                                                                                Query Match
Best Local :
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                                                                                                                                                                                                                                                                                                                                                                                                             ORGANISM: Artificial Sequence FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: PRT
                                                                                                                                                                                               32 QVYYRPVDRYSNQ 44
                                                                                                                                                                                                                                                                        Local Similarity 69.
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                                                                                                                                                                                                                                                                                          87.98;
69.28;
    IMMUNOLOGICAL ASSAY FOR SPONGIFORM ENCEPHALOPATHIES
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Pred. No. 0.0016;
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NUMBER OF SEQUENCES: 4
COMPUTER READABLE FORM:

MEDIUM TYPE:

Floppy disk

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RESULT 4
US-09-745-003-9
; Sequence 9, Application US/09745003
; Patent No. US20020042122A1
. GENERAL INFORMATION:
                                                                                                                                                                                                                                                      ; LENGTH: 161
; TYPE: PRT
; ORGANISM: sheep
US-09-745-003-7
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Patent No. US20020042122A1
GENERAL INFORMATION:
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                                                                                                                                                                                                                                                                                                                    SEQ ID NO 7
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                                                                                                                                                                                             Matches
                                                                                                                                                                                                                        Query Match
                                                                                                                                                                                                                                                                                                                               TITLE OF INVENTION: Human Proteins; Related Reagents FILE REFERENCE: PPP2 CURRENT APPLICATION NUMBER: US/09/745,003 CURRENT FILING DATE: 2000-12-20 NUMBER OF SEQ ID NOS: 13 SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Bazan, Fernando J
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEFAX: 353-1-660692
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30 (EPO)
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ORIGINAL SOURCE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: amino acid HYPOTHETICAL:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
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APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ANTI-SENSE:
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nes 9; Conserv
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CELL TYPE:
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                                                                                                                                  68 QVYYRPVDRYSNQ 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: unknown
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                                                                                                                                                              1 QVYYXPXDXYXNQ 13
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es 9; Conservative
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TD NO: 4:
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69.2%;
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69.2%;
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Pred. No.
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Pred. No.
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QVYYRPVDQYSNQ 81

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Best Local Similarity
watches 9; Conserve
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; ORGANISM: bovine
US-09-745-003-9
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US-09-745-003-10
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                                                                                              ; ORGANISM: rodent
US-09-745-003-12
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CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: PATENTIN VEr. 2.0
SEQ ID NO 9
LENGTH: 161
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/745,003
CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 10
LENGTH: 162
                                                                                                                                                      CURRENT APPLICATION NUMBER: US/09/745,003
CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 12
SEQ ID NO 12
                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: Bazan, Fernando J
TITLE OF INVENTION: Human Proteins; Related Reagents
                                                                                                                                                                                                                                                                                                 Sequence 12, Application US/09745003 Patent No. US20020042122A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               GENERAL INFORMATION:
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Patent No. US20020042122A1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
                                                            Query Match
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                                Best Local Similarity Matches 9; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: Human Proteins; Related Reagents FILE REFERENCE: PrP2
                                                                                                                                                                                                                                          FILE REFERENCE: PrP2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Bazan, Fernando J
TITLE OF INVENTION: Human Proteins; Related Reagents
                                                                                                                               TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: PRT
                                                                                                                                            LENGTH:
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 1 QVYYXPXDXYXNQ 13
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                                  Conservative
                                               87.9%;
69.2%;
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69.2%;
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                                             Score 51;
Pred. No.
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Pred. No. 0.
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                                  Mismatches
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0.0055;
                                                 0.0056;
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                                                                 DB 10;
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LENGTH: 253
; TYPE: PT
; ORGANISM: homo sapiens
; PUBLICATION INFORMATION:
; DATABASE EACCESSION NUMBER: NCBI ENTREZ / XM_009567
; DATABASE ENTRY DATE: 2001-04-17
; RELEVANT RESIDUES: (1)..(253)
US-09-904-987-3
                                                                             RESULT 9
US-09-943-906-2
                                                                                                                                                                                                                                                                                                                  : NAME/KEY: misc_feature
; OTHER INFORMATION: Incyte ID No. US20020119463A1 1256895CD1
US-09-919-172-57
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      US-09-919-172-57
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                  Sequence 2, Application US/09943906 Patent No. US20020150571A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 3, Application US/09904987 Patent No. US20020037908A1 GENERAL INFORMATION: GENERAL INFORMATION: APPLICANT: No. US20020037908A1actyl, Inc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           GENERAL INFORMATION
                                                                                                                                                                                                                                                                                   Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                             SOFTWARE: PERL Program
SEQ ID NO 57
LENGTH: 253
TYPE: PRT
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Turner, Christopher M.
TITLE OF INVENTION: PROSTATE CANCER MARKERS
FILE REFERENCE: PA-0036 US
CURRENT APPLICATION NUMBER: US/09/919,172
CURRENT FILING DATE: 2001-07-30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIOR APPLICATION NUMBER: 60/222,469
PRIOR FILING DATE: 2000-07-29
NUMBER OF THE PRIOR FILING DATE: 2000-07-29
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CURRENT FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Faris, Mary APPLICANT: Turner, Ch.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TITLE OF INVENTION: Methods and Compositions for Controlling Pathological and Prepath TITLE OF INVENTION: Protein Assembly or Aggregation FILE REFERENCE: 42108/26146
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: PatentIn version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NUMBER OF SEQ ID NOS: 102
                                                                                                                                                                                                                                                                                                                                                                                                      ORGANISM: Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                       FEATURE:
                                                                                                                                                               160 OVYYRPMDEYSNO 172
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                                                                                                                                                                                                                                           Local Similarity
nes 9; Conserv
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    APPLICANT:
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ilarity 69.2%;
Conservative
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Prusiner, Stanley B
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Pred. No. 0.0088;
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Pred. No. 0.0088;
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                                                                                                                                                                                                                                                                                     DB 10;
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RESULT 10
US-10-106-574-5
Query Match
Best Local Similarity
Conserva
                                                                                    ; LENGTH: 254
TYPE: PRT
; ORGANISM: Murinae gen. sp.
US-10-106-574-5
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                                                                                                                                                       CURRENT APPLICATION NUMBER: CURRENT FILING DATE: 2002-C NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn version 3
SEQ ID NO 5
                                                                                                                                                                                                                                                                                                                     Sequence 5, Application US/10106574 Patent No. US20020164335A1 GENERAL INFORMATION:
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Matches 9; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
                                                                                                                                                                                                                                             APPLICANT: Harris, David A.
APPLICANT: Stewart, Richard S.
TITLE OF INVENTION: Compositions and Methods for the Study and Diagnosis of Prion FILE REFERENCE: 09789280.0003
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                160 QVYYRPMDEYSNQ 172
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APPLICATION NUMBER: 09/550,374
FILING DATE: <UNKNOWN>
ATTORNEY/AGENT INFORMATION:
NAME: BOZICEVIC, Karl
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
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MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TOPOLOGY: linear MOLECULE TYPE: peptide
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
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Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICATION NUMBER: US/09/943,906
FILING DATE: 30-Aug-2001
CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TYPE: amino acid
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                  Conservative
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                                                                                                                                                                           version 3.1
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                                   87.9%;
69.2%;
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69.28;
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                                   Score 51; DB 9;
Pred. No. 0.0088;
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Pred. No. 0.0088;
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RESULT 13
US-10-106-574-8
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US-10-106-574-6
                                        GENERAL INFORMATION:
APPLICANT: Harris, David A.
APPLICANT: Stewart, Richard S.
TITLE OF INVENTION: Compositions and Methods for the Study and Diagnosis of Prion
FILE REFERENCE: 09789280.0003
FILE REFERENCE: 09789280.0003
                                                                                                                                              Sequence 8, Application US/10106574 Patent No. US20020164335A1
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             CURRENT APPLICATION NUMBER: US/10/106,574
CURRENT FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION NUMBER: US/10/106,574
CURRENT FILING DATE: 2002-03-26
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TITLE OF INVENTION: Compositions and Methods for the Study and Diagnosis of Prion Dis
FILE REFERENCE: 0978930.0003
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SOFTWARE: PatentIn version 3.1
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CURRENT FILING DATE: 2002-03-26
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69.2%;
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Best Local Similarity
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; TYPE: PRT
; ORGANISM: Murinae
US-10-106-574-8
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                                                                                                                                                                                                                                TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 1:
SEQUENCE CHARACTERISTICS:
159 QVYYRPVDQYSNQ 171
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                            1 QVYYXPXDXYXNQ 13
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SOFTWARE: FastSEQ Version 2.0
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                          TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                      MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                  PRIOR APPLICATION DATA:
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ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NUMBER OF SEQUENCES: 86
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Prusiner, Stanley B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP
                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/943,906 FILING DATE: 30-Aug-2001
                                                                                                                                                                                                TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                       NAME: Bozicevic, Karl
REGISTRATION NUMBER: 28,807
                                                                                                                                                                                                                                                                                                                                                                                       APPLICATION NUMBER: 09/550,374 FILING DATE: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                  CLASSIFICATION: <Unknown>
                                                                                                                                                                   TOPOLOGY: linear
                                                                                                                                                                                   STRANDEDNESS: single
                                                                                                                                                                                                                   LENGTH: 254 amino acids
                                                                                                                                                                                                                                                                             TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COMPUTER: IBM Compatible
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                                                           Conservative
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Burton, Dennis R.
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                                                         Score 51; DB 10;
Pred. No. 0.0088;
0; Mismatches 4
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Pred. No. 0.
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RESULT 15

US-09-943-906-4

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PRICE A. Application US/09943906

Patent No. US202015971A1

PRICENT US202015971A1

PRICENT US202015971A1

APPLICANT: PIVALER, Stanley B.

HILLAMSON, R. ANTHONY

HILLAMSON, R. ANTHONY

BUTCOP INVENTION: ANTHONY

BUTCOP FOR NATIVE PTP

NUMBER OF SEQUENCES: 86

CORRESPONDENCE ADDRESS:

ADDRESSES: 1816 ARTHONY

FREET: 200 Sand Hill Road

CITY: Menilo Park

COUNTRY: U. S. A.

COUNTRY: U. S. A.

COUNTRY: U. S. A.

COMPUTER REDARBLE FORM:

MEDIUM TYPE: Diskette

COMPUTER: SIDABLE FORM:

MEDIUM TYPE: DISKETTE

MOLECULOR: CHARACTERISTICS:

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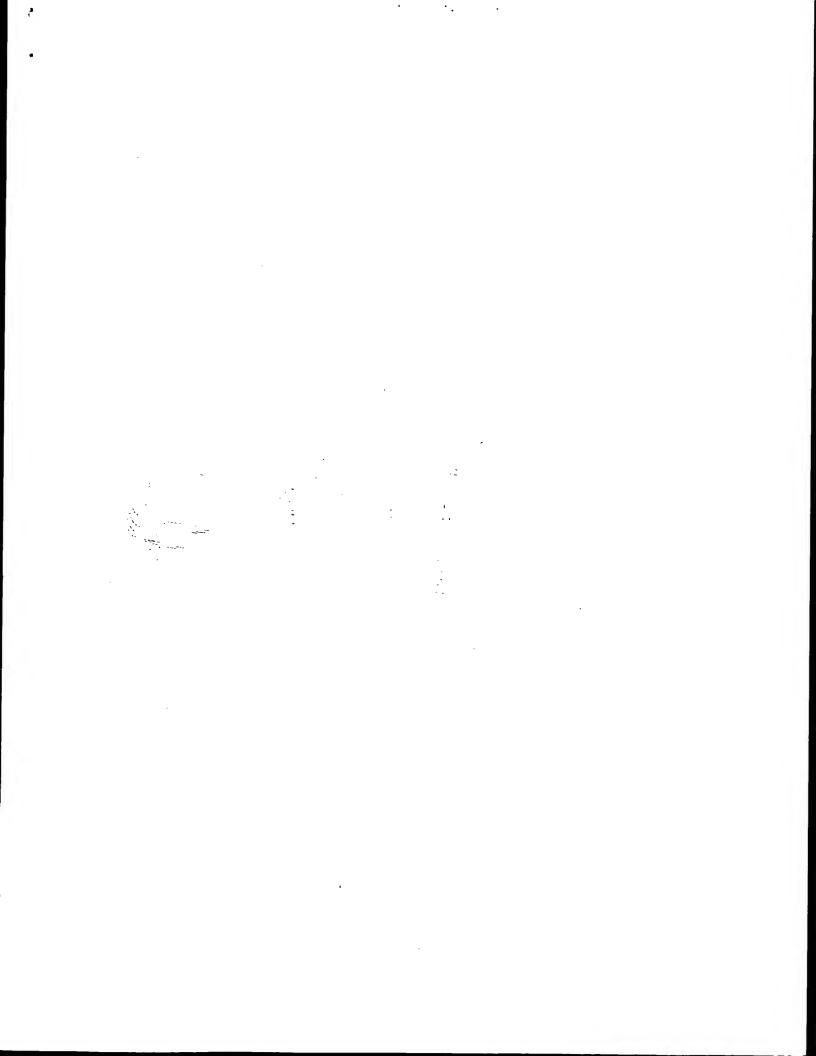
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MOLECULOR: CHARACTERISTICS:

MOLECULOR: TYPE: DISKETTION: SEQUENCE SIDABLE FORM:

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Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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Maximum DB seq length: 2000000000
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

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41	42	42	42	42	42	42	42	42	42	Score	
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АВВ08378	ABB04428	AAE15608	AAB61773	AAB82117	AAP93673	AAB72369	AAB30802	AAW92807	AAW17678	ID	
Mutant immunogenic	Hamster prion prot	Hamster PrP protei	Hamster prion prot	Hamster Prp. Cric	Partial nucleotide	Rabbit prion prote	Amino acid sequenc	Mouse rPrP protein	Syrian hamster pri	Description	

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ALIGNMENTS

RESULT 1 AAW17678

AAW17678 standard; peptide; 141 AA.

14-JAN-1998 (first entry)

Syrian hamster prion protein peptide SHa 90-231.

Prion protein; PrP; alpha helical domain; screening; inhibition; binding; scrapie; bovine spongiform encephalopathy; BSE; CJD; Creutzfeldt-Jakob disease; kuru; GSS; FFT; fatal familial insomnia; Gerstmann-Straussler-Scheinker disease; hamster; human.

Mesocricetus auratus.

W09716728-A1.

09-MAY-1997.

28-OCT-1996; 96WO-US17462.

02-NOV-1995; 95US-0556823.

(REGC) UNIV CALIFORNIA.

Cohen FE, Kaneko K, Prusiner SB;

WPI; 1997-272248/24.

Prion proteins (PrPs) having at least one alpha-helical domain - used in assays for screening compounds able to inhibit or decrease

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RESULT 2
AAW92807
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                        This invention describes a composition of Prion Protein Modulation Fractor (PPMF) which binds the prion protein PrP(c) and facilitates a conformational change from PrP(c) to the scrapic isoform of the PrP protein, PrP(sc). PPMF is involved in complex formation as the rate limiting step. The protein can therefore be used in assays to 'speed up', formation of the complex and conversion of prion proteins to the pathogenic stage and thus can used to rapidly detect the presence of pathogenic prion proteins in a sample. The compositions can be used to screen for compounds that inhibit PrP(sc) formation. The gene can also be used to generate transgenic animals which are
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Murine; prion protein; PPMF; prion protein modulator factor; PrP; PrP(c); PrP(sc); scrapie; isoform; pathogenic; inhibitor; screening; disease resistance; transgenic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present sequence represents a prion protein (PrP) peptide from syrian hamster. PrP has an ability to induce a conformational change in cellular prion protein (PrP-c). Methods, for screening compounds which inhibit the binding of PrP-c to a PrP peptide, are used for screening for drugs that may be useful in the treatment prion-related disease e.g. scraple, BSE (bowine spongiform encephalopathy). CJD (Creutzfeldt Jakob disease), kuru, GSS (Gerstmann-Straussler-Scheinker
                                                                                                                                                                                                                                                                                                                                                                                                       12-MAY-1998;
02-JUN-1997;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             07-MAY-1999
                                                                                                                                                                                                                   Disclosure; Fig 6A-B; 93pp; English.
                                                                                                                                                                                                                                                 Novel Prion Protein Modulation Factor(s) - useful to increase speed and sensitivity of assays to detect pathogenic prion proteins
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               resistant to prion diseases.
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les 8; Conserv
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97US-0868162
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Pred. No. 0.07
0; Mismatches
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CTTQYQKESQAYY 206

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RESULT 3
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                                                                                                              The present sequence represents a prion protein. The specification codescribes chimeric polypeptides, which comprise at least one SCHAG (self-coalesces into higher-order aggregates) amino acid sequence fused in frame with a polypeptide of interest (which is other than a marker protein, a glutathione-S-transferase or a staphylococcal nuclear protein). The specification also describes chimeric polypeptides that comprises an amyloidogenic domain that causes aggregation into fibrils. The chimeric polypeptides are used to prepare polymers with multiple reactivities, e.g. derivatised with enzymes, or specific binding partners, and useful e.g. for performing multi-step chemical reactions. They can be used create an inducible, or stable phenotypic alteration in a cell, e.g. for gene therapy, protein production, imparting disease resistance to plants, altering plant pigmentation and for diagnosis cand treatment of prion diseases.
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Best Local (
                                 Matches
                                          Query Match
Best Local
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                                                                                                                                                                                                                                                                                                                                         Claim 11; Page 139-140; 188pp; English.
                                                                                                                                                                                                                                                                                                                                                                 New nucleic acid encoding chimeric proteins with self-assembly properties, useful e.g. for diagnosis and treatment of prion diseases, also related aggregates, fibrils and polymers -
                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-061723/07.
N-PSDB; AAC86687.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SCHAG; self-coalesce; higher-order aggregate; amyloidogenic domain;
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                                                                                        Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            aggregation; fibril; phenotypic alteration;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Amino acid sequence of a Syrian hamster prion protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   125 CTTQYQKESQAYY 137
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 1 CXTQYXXESXAXY 13
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8; Conserv
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                                            82.4%;
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                                            Pred. No. 0.
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Pred. No. 0.077;
                                 Mismatches
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RESULT 5
AAP93673
ID AAP9
XX
AC AAP9
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                                                                                                                                                                                                                                                               This invention relates to a peptide fragment of a cellular form of prion protein PrPC located around a disulphide bond found in PrPC. The stable structure is a specific marker of PrPC but not soluble prion protein (PrPSC). The PrPC peptide sequences can be used to generate an antibody or binding agent that binds PrPC. The antibody is used to detect or remove PrPC, and may be used in preventative medicine. The antibody may be used in the prevention, treatment or diagnosis of a prion disease, e.g. spongiform encephalopathies, such as Scrapie in sheep, bovine spongiform encephalopathies (BSE) in cows, and Creutzfeldt-Jakob disease (CJD) in humans. The present sequence represents the cellular form of rabbit prion protein, the stable region of the protein may be used in the production of anti-PrPC antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Key
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 New prion peptide for treating, preventing and/or diagnosing prion diseases e.g. scrapie in sheep, bovine spongiform encephalopathies cows and Creutzfeldt-Jakob disease in humans
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30-JUL-1999;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   17-MAY-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             AAB72369;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB72369 standard; Peptide; 250 AA.
  31-MAY-1990
                                                                                                                                                                                                                                           Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 3; Fig 5; 69pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25-JUL-2000; 2000WO-GB02873
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (IMCO-) IMPERIAL COLLEGE INNOVATIONS LTD.
                             AAP93673;
                                                      AAP93673 standard;
                                                                                                                             211 CITQYQQESQAAY 223
                                                                                                                                                      1 CXTQYXXESXAXY 13
                                                                                                                                                                                                                                         250 AA;
                                                                                                                                                                                  Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (first entry)
  (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               99GB-0017491.
99GB-0017878.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Location/Qualifiers
176..221
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             /note=
                                                      protein; 254
                                                                                                                                                                                                82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Stable region, specifically claimed in claim
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Waltho JP, Jackson GS,
                                                                                                                                                                                  0;
                                                                                                                                                                                                 Score 42; DB 2
Pred. No. 0.14;
                                                                                                                                                                                     Mismatches
                                                                                                                                                                                                               22;
                                                                                                                                                                                    5,
                                                                                                                                                                                                             Length 250;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Hosszu
                                                                                                                                                                                     Indels
                                                                                                                                                                                  0,
                                                                                                                                                                                    Gaps
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RESULT 6
AAB82117
ID AAB8
XX AAB82
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XX OS Cric
XX Us62
XX Us62
XX Us63
XX Us63
XX PF 03-A
XX PF 03-A
XX Us8
XX Uses
PA (Us8
XX Uses
PA (Us8
                                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
Best Local Similarity
Watches 8; Conserve
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Detecting susceptibility to scrapie in sheep, cattle by analysing blood or tissue for polymorphism linked susceptibility, pref. using DNA hybridisation probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          17-MAY-1988;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W08911545-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cricetulus sp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        scrapie associated
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Scrapie; PrP gene;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Partial nucleotide sequence of Hamster PrP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Product of partial hamster PrP gene, which is sufficiently similar the sheep Sip gene (scrapie susceptibility), that the probe (pEA974 be made to the site.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Disclosure; ; 46pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   N-PSDB; AAN92724.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Hope J, Hunter N;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 15-MAY-1989;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            30-NOV-1989
                                                                                                                                                                            Hamster; Prp; cerebroprotective; Prp conversion inhibitor; prion protein; transmissible spongiform encephalorathy; TSE; neurodegenerative disease; protease-sensitive prion protein; PrPsen; protease-resistant prion protein; PrPres.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1989-370736/50.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (ANIM-) ANIMAL HEALTH LTD.
                                                 03-AUG-1998;
                                                                          03-AUG-1998;
                                                                                                     03-APR-2001
                                                                                                                            US6211149-B1
                                                                                                                                                     Cricetulus griseus.
                                                                                                                                                                                                                                           Hamster Prp.
                                                                                                                                                                                                                                                                     29-JUN-2001
                                                                                                                                                                                                                                                                                                AAB82117;
                                                                                                                                                                                                                                                                                                                        AAB82117 standard; Protein; 254 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      See also AAN92735.
                                                                                                                                                                                                                                                                                                                                                                                         214 CTTQYQKESQAYY 226
                                                                                                                                                                                                                                                                                                                                                                                                                 1 CXTQYXXESXAXY 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              254 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         88GB-0011608
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   89WO-GB00522.
                                                 98US-0128450.
                                                                           98US-0128450.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Bovine spongiform encephalopathy; BSE; fibrils; SAF.
                                                                                                                                                                                                                                                                                                                                                                                                                                                       82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 42; DB 10;
Pred. No. 0.14;
                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                           5
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 254;
                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        and
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                                                                                                                                                                                                                                                                                                                                                                                                                                           0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (pEA974) can
                                                                                                                                                                                                                                                                                                                                                                                                                                           Gaps
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(USSH) US DEPT HEALTH & HUMAN SERVICES.

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Caughey BW,

Chabry J,

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local
                                                                   New antibody for diagnosing, preventing and treating such as Creutzfeldt-Jakob disease, Kuru, insomnia in
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Transmissible spongiform encephalopathies (TSE) are fatal neurodegenerative diseases. These diseases are characterised by the formation and accumulation, in the brain, of an abnormal proteinase K realstant isoform (Prpres) of a normal protease-sensitive host-encoded prion protein (Prpsen). The present invention relates to peptides comprising a hamster, human or murine prion protein (Prp) fragment which specifically inhibit the conversion of protease-sensitive prion protein (Prpsen) to protease-resistant prion protein (Prpres). The present sequence is the protein sequence for hamster Prp. The peptides of the present invention are useful for diagnosis and treatment of TSE
                         Disclosure; Fig 2; 81pp; English.
                                                                                                                   WPI; 2001-102614/11.
                                                                                                                                               Pinard
                                                                                                                                                              Cashman NR, Paramithiotis E,
                                                                                                                                                                                                                                                                                                                 WO200078344-A1
                                                                                                                                                                                                                                                                                                                                                                                                    Prion; neuroprotective; vaccine; prion protein peptide; PrP; PrPSC; immunisation; antigenic; antibody; Creutzfeldt-Jakob disease; CJD;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             New peptide comprises hamster prion protein fragment that inhibits conversion of prion protein from protease-sensitive to protease-resistant form, useful for diagnosis and treatment of spongiform encephalopathies
                                                                                                                                                                                                                                                     23-JUN-2000; 2000WO-US17455
                                                                                                                                                                                                                                                                                   28-DEC-2000.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Hamster prion protein (PrP) sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                20-APR-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              AAB61773;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         AAB61773 standard; protein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence
                                                                                                                                                                                           (CAPR-) CAPRION PHARM INC.
                                                                                                                                                                                                                                                                                                                                             Mesocricetus auratus
                                                                                                                                                                                                                                                                                                                                                                          spongiform
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Disclosure; Column 39-42; 31pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        N-PSDB;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                214 CTTQYQKESQAYY 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity es 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CXTOYXXESXAXY 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        2001-315407/33.
DB; AAF86478.
                                                                                                                                                                                                                                                                                                                                                                   Gerstmann-Straussler syndrome; fatal familial insomnia; scrapie; lform encephalopathy; hamster.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      254 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (first entry)
                                                                                                                                                                                                                       99US-0140634.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        82.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          254 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Score 42;
Pred. No.
                                                                                                                                                            Slon-Usakiewicz J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0.14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 22;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Length 254;
                                                                                                                                                            Haghighat A;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0;
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The invention relates to an antibody (I) that binds with high-binding

The present invention relates to peptides comprising a peptide region protein (PrP) and which are capable of inhibiting conversion of protease sensitive prion protein (PrPsen) to protease resistant prion protein (PrPres). The peptides are useful for inhibiting formation of protease resistant prion proteins, such as those associated with

of.

transmissible spongiform encephalopathies, e.g..

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AAE15608
ID AAE1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    affinity to a YYX epitope of a mammalian prion protein peptide (PrP),

PrPSC. (I) is useful for detecting PrPSC in a biological sample such as

can tissue or cell, their extracts, a bodily fluid or a biopsy from a

chaman, livestock species, or a pet species, by forming a complex between

prPSC and (I). (I) is useful for preventing or treating a PrPSC disease

cin a mammal, for decontaminating PrPSC from a biological sample or for

cinhibiting PrPSC in a biological sample, where the sample is perfused

cc with the antibody. A vaccine comprising (I) or a peptide that has

cc antigenicity as a PrPSC is useful for immunizing a mammal against PrPSC

cdisease. Such antigenic peptides and synthetic peptides are useful as

cc immunogens for generating antibodies specific for PrPSC. The prion

cc disease such antigenic peptides and synthetic peptides are useful as

cc immunogens for generating antibodies specific for PrPSC. The prion

cc disease (CJD), variant CJD, iatrogenic CJD, familial CJD, Kuru, Gerstmann

cc disease (CJD), variant CJD, iatrogenic CJD, familial CJD, Kuru, Gerstmann

cc sheep and goats and spongiform encephalopathy in cattle. The present

cc sequence represents a hamster prion protein amino acid sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local S
Matches 8
                                                                                                                                                           Peptides which can inhibit conversion of prion proteins from protease sensitive to protease resistant forms, are useful in treating
                                                                                                                                                 sensitive to protease resistant forms, ar transmissible spongiform encephalopathies
                                                                                                                                                                                                                                                                       Chesebro BW,
                                                                                                                                                                                                                                                                                                                                           12-MAY-1998;
03-AUG-1998;
                                                                                                                                                                                                                                                                                                                                                                                           30-MAR-2001; 2001US-0823494.
                                                                                                                                                                                                                                                                                                                                                                                                                                                               US2001041790-A1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cricetulus sp.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gerstmann-Straussler-Scheinker syndrome; scrapie; kuru; hamster.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Protease resistant prion protein; PrPres; Creutzfeldt-Jakob disease; protease sensitive prion protein; PrPsen; therapy; neuroprotectant; transmissible spongiform encephalopathy; fatal familial insomnia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        AAE15608;
                                                                                                                                                                                                                                                                                                         (USSH ) US DEPT HEALTH & HUMAN SERVICES.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hamster PrP protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAE15608 standard; Protein; 254 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                               15-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      12-MAR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            214 CTTQYQKESQAYY 226
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Local Similarity es 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 CXTQYXXESXAXY 13
                                                                                                                                                                                                                                     2002-065984/09.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                254 AA;
                                                                                                                Page
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                                                                                                                                                                                                                                                                         98US-085160P.
98US-0128450.
                                                                                                             22; 33pp;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             82.4%;
                                                                                                                                                                                                                                                                       BW,
                                                                                                              English.
                                                                                                                                                                                                                                                                   Chabry J, Priola
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 42; L
Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             BG
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . 14;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 254;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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33333333<del>3</del>3
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Best Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              disease, kuru, Gerstmann-Straussler-Scheinker syndrome, fatal familial insomnia or scrapie. The peptides can be used as diagnostic agents, e.o. to detect the presence of PrPres in body fluids such as blood. They may also be used to treat or prevent diseases such as the above. They may also be used to design analogues, derivatives or mimetics for use as inhibitors of conversion of PrPsen to Prpres. The present sequence is
                                                                                                  The present invention relates to normal (PrPc) or abnormal (PrPsc) human or animal prion proteins which are able to bind to DNA or RNA, particularly of viral, especially retroviral, origin and to nucleocapsid proteins (NCP) of human or animal retroviruses. These can be used as antiviral agents, particularly against human immune deficiency virus (HIV), and in the treatment of prion diseases including Creutzfeld-Jakob disease, Gerstmann-Straeussler-Scheinker disease, fatal familial insommia, kuru, bovine spongiform encephalitis and scrapie. The present sequence is the hamster PrP protein sequence.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Cricetulus griseus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Hamster; prion protein; PrP; antiviral; HIV; prion disease; kuru; antibacterial; neuroprotective; anti-HIV; Creutzfeld-Jakob disease; Gerstmann-Straeussler-Scheinker disease; fatal familial insomnia; bovine spongiform encephalitis; scrapie; virucide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABB04428 standard; Protein; 257
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                             Disclosure; Fig 9; 80pp; French.
                                                                                                                                                                                                                                                                                                            New polypeptides, useful as antiviral agents, comprise their prion proteins able to bind nucleic acid, nucleocapsid proteins, and ligands for use as antiprion agents -
                                                                                                                                                                                                                                                                                                                                                                                WPI; 2002-049350/06.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            WO200183747-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Hamster prion protein PrP.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ABB04428;
                                                                      Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                    Leblanc P,
                                                                                                                                                                                                                                                                                                                                                                                                                                                   (INRM ) INSERM INST NAT SANTE & RECH MEDICALE.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     28-APR-2000; 2000FR-0005535.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       30-APR-2001; 2001WO-FR01336.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         08-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  hamster PrP protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          214 CTTQYQKESQAYY 226
Local Similarity
hes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Local Similarity
mes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CXTQYXXESXAXY 13
                                                                       257 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 254 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                   Darlix J,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               82.4%;
61.5%;
                 82.4%;
61.5%;
                                                                                                                                                                                                                                                                                                                                                                                                                    Gabus-Darlix C;
   0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Score 42; DB 2
Pred. No. 0.14;
 Score 42; DB 2:
Pred. No. 0.14;
); Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mismatches
                                  DB 23; Length 257;
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1 CXTQYXXESXAXY 13

Query Match
Best Local Similarity
Themes 8; Conserv

Conservative

80.4%;

Score 41; [Pred. No. 0. Mismatches

DB 23; .01;

Length 13; Indels

0

Gaps

0

Sequence

13

A,

antibodies specifically against PrPSc.

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ABB08378
ABB08378
AC ABB0
XX ABB0
XX ABB0
DT 22-A
DT 22-A
DX WITH
XX W HITH
KW CJD;
XX W TTAI
XX EP1:
XX EP1:
XX EP1:
XX W TTAI
XX W TTA
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                                                                                                              The invention relates to an antibody selectively binding to the three cdimensional conformation of the C-terminal of the infectious 'scraple' (C (PrPSC) isoform of the prion protein (or a part), but not binding to the structure of the C-terminal of the normal cellular (PrPC) isoform of the grion protein. The antibodies (or functional parts) are useful in the C diagnosis of Bovine Spongiform Encephalopathy (BSE), Creutzfeldt Jacobs (C Spongiform Encephalopathy (TSE) related diseases e.g. In humans, cows, (C sheep etc., since they can differentiate between the infectious (PrPSC) isoform and the normal cellular (PrPC) isoform of the prion protein.

C They are also useful in the treatment of such diseases. They may be included in pharmaceutical compositions or used to produce drugs or cative and/or passive vaccines against diseases as above. The antidiotypic antibodies of the polypeptides of the invention may also be used to produce drugs or vaccines against such diseases. The current sequence represents a mutant immunogenic peptide derived from bovine c prion protein C-terminal region (see ABB08377). This sequence is created c from the wild-type sequence by replacement of the Gln residue at position and the peptide with Glu. This sequence has proven to elicit an immune cannot be produced to the peptide with Glu. This sequence has proven to elicit an immune cannot be produced to the peptide and the produce and the position control the call the produce and the position and the produce and the produce and the position control the call the produce and the position cannot be produced to be capable of easily raising
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        New antibody, useful in diagnosis and treatment of BSE, CJD, new variant CJD and other Transmissible Spongiform Encephalopathy related diseases, selectively binds to the infectious form of the prion protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Prion; BSE; bovine spongiform encephalopathy; vCJD; new variant creutzfeldt jacobs disease; scrapie; TSE; transmissible spongiform encephalopathy; antibody; PrPSc; PrPC; vaccine;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ABB08378 standard; peptide; 13 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; Page 10; 21pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  23-MAY-2000; 2000EP-0111108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         23-MAY-2000; 2000EP-0111108.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Mutant immunogenic peptide derived from bovine prion protein.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  22-APR-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ABB08378;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          (BLOO-) BLOOD TRANSFUSION CENT SLOVENIA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              217 CTTQYQKESQAYY 229
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                2002-107827/15
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            creutzfeldt jacobs disease; cow.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Location/Qualifiers
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CXTQYXXESXAXY

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ABB08379
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                                                                                                                                                                                The invention relates to an antibody selectively binding to the three dimensional conformation of the C-terminal of the infectious 'scrapie' (PrPSc) isoform of the prion protein (or a part), but not binding to the Structure of the C-terminal of the normal cellular (PrPc) isoform of the prion protein of the normal cellular (PrPc) isoform of the prion protein. The antibodies (or functional parts) are useful in the diagnosis of Bovine Spongiform Encephalopathy (BSE), Creutzfeldt Jacobs Disease (CJD), new variant form CJD (VCJD) and other Transmissible Spongiform Encephalopathy (TSE) related diseases e.g. in humans, cows, is the prior protein. The stream and the normal cellular (PrPc) isoform of the prion protein. They are also useful in the treatment of such diseases. They may be included in pharmaceutical compositions or used to produce drugs or cantidotypic antibodies of the polypeptides of the invention may also be used to produce drugs or vaccines against diseases. The current sequence represents an immunogenic peptide derived from bovine prion cover.
                                                                             Matches
                                                                                                                 Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  New antibody, useful in diagnosis and treatment of BSE, CJD, new variant CJD and other Transmissible Spongiform Encephalopathy related diseases, selectively binds to the infectious form of the prion protein
                                                                                                                                                        Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Curin-Serbec V;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                new variant creutzfeldt jacobs disease; scraple; TSE; transmissible spongiform encephalopathy; antibody; PrPSc; PrPC; vaccine; CJD; creutzfeldt jacobs disease; cow.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prion; BSE; bovine spongiform encephalopathy; vCJD; new variant creutzfeldt jacobs disease; scrapie; TSE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 18;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     23-MAY-2000; 2000EP-0111108
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (BLOO-) BLOOD TRANSFUSION CENT SLOVENIA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          23-MAY-2000; 2000EP-0111108
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  28-NOV-2001.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Bos taurus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          22-APR-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ABB08379;
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                                                                           Local Similarity es 8; Conserv
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CITQYQRESQAYY 13
                                    CXTQYXXESXAXY 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Page 11; 21pp; English.
                                                                                                                                                        13 AA;
                                                                           Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                peptide derived from bovine prion protein.
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                                                                                           80.4%;
                                                                           0;
                                                                                         Score 41; [
Pred. No. 0.
                                                                         Mismatches
                                                                                         DB 2:
                                                                                                           23;
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                                                                                                             Length 13;
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                                                                     Gaps
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RESULT 13
AAB69083
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AAR38048
                                                                                                                                                                                                                  Query Match
Best Local
                                                                                                                                                                                                        Matches
                                                                                                                                                                                                                                                                                                   The sequences given in AAR38041-48 represent polypeptides which are derived from an antigenic site, region F, of a prion protein. Prion proteins comprise six regions of interest (A-F), and two related frame shift peptides sequences caused by a repeating section in region E having a nucleic acid coding sequence frame shift mutation of +1 (FSa) or -1 (FSb). These peptides (see also AAR38041-48) and antibodies raised against these may be used to treat or prevent spongiform encephalopathy in humans, sheep or cattle. They can be used to block ceilular binding and aggregation of prion proteins and to stimulate the mammalian immune system. These peptides may be used to distinguish between the normal form of prion protein (PrPc) and the scrapic-associated form (PrPsc). These peptides may include rare or synthetic amino acids or a ratio-inverso peptide modification to improv
                                        AAB69083;
                                                                  AAB69083 standard;
                                                                                                                                                                                                                                                                 Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Claim 29;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               New polypeptide(s) contg. antigenic site of prion protein - useful for treatment and diagnosis of mammalian encephalopathies e.g. Creutzfeld-Jacob disease and kuru
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Fishleigh RV,
                                                                                                                                                                                                                                                                                             resistance to enzymatic degradation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             WPI; 1993-196994/24.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  03-DEC-1991;
10-JUL-1992;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               03-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             10-JUN-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Antigen; prion; protein; region; frame shift; repeat; mutation; PrPc; FSa; FSb; subfragment; antibody; treatment; spongiform encephalopathy; human; sheep; cattle; cellular binding; aggregation; mammal; scrapie; immune system; PrPsc; ratio-inverso peptide; enzymatic degradation;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prion protein region F #2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       14-OCT-1993
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Synthetic.
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                                                                                                                                                                         1 CXTQYXXESXAXY 13
                                                                                                                                           CITOYORESOAYY 13
                                                                                                                                                                                                        Similarity
8; Conser
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Page 74; 82pp; English.
                                                                                                                                                                                                                                                               15 AA;
                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Mee RP,
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92GB-0014663.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               92WO-GB02246.
                                                                  peptide; 17 AA
                                                                                                                                                                                                                      80.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Robson B;
                                                                                                                                                                                                      0;
                                                                                                                                                                                                                  Score 41; DB 14; Length 15; Pred. No. 0.012;
                                                                                                                                                                                                        Mismatches
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                                                                                                                                                                                                     Gaps
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23-APR-2001

(first entry)

Bovine

prion protein peptide SEQ ID NO:3.

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                                                                                                                                                                                                                                                                                                                                                                                                                       Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  specifically binds to a conserved epitope of prion proteins, which is capable of binding prion protein (PrP)-Sc protein in fixed or unfixed tissue that has been treated to unmask the group to PrP-Sc protein and eliminate availability of a corresponding group of PrP-Cellular. (I) can be used in immunoassays to detect PrP-Sc proteins in animal or human, including ruminant livestock, cats, mink and non-human primates, sheep, goat, cattle, mule deer and elk, as an indication of the presence of transmissible spongiform encephalopathies (TSE). Presence of PrP-Sc indicates the scrapic, bovine encephalopathy- or chronic wasting disease-infected animals. The antibodies provide early detection of PrP-Sc and for preclinical diagnosis of TSE. The present sequence represents a prion protein peptide which is used in an example from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Monoclonal antibody; detection; prion protein; TSE; infection; transmissible spongiform encephalopathy; scrapie; bovine encephalopathy; chronic wasting disease; PrP-Sc.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Monoclonal antibodies for detecting prior proteins as an indication of transmissible spongiform encephalopathies, specifically binds to conserved group of prior proteins -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  15-JUL-1999;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                14-JUL-2000; 2000WO-US19211.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   25-JAN-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Bos taurus
                                           transmissible spongiform encephalopathy; prion protein; sterilisation; immunisation; Creutzfeldt-Jacob disease; kuru; fatal familial insomnia; Gerstmann-Straussler-Scheinker syndrome; chronic wasting disease; bovine spongiform encephalopathy; feline spongiform encephalopathy; scrapie; transmissible mink encephalopathy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The present invention describes a monoclonal antibody (I) which
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Example 2; Page 12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (USDA ) US DEPT OF AGRICULTURE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    WO200105426-A1
                                                                                                                                                                Prion mimetic peptide SEQ ID NO:4.
                                                                                                                                                                                                                                                                  ABB81632 standard; peptide; 19 AA
             Synthetic
                                                                                                                                 Prion mimetic
                                                                                                                                                                                                25-SEP-2002 (first entry)
                                                                                                                                                                                                                                     ABB81632
                                                                                                                                                                                                                                                                                                                                                                                                                                       Local
                                                                                                                                                                                                                                                                                                                                                                                      1 CXTQYXXESXAXY 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           present invention.
                                                                                                                                                                                                                                                                                                                                               CITOYORESOAYY 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              2001-159477/16.
                                                                                                                                                                                                                                                                                                                                                                                                                     Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           17
                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3
                                                                                                                             peptide; degradation; detection;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  9908-0353348.
                                                                                                                                                                                                                                                                                                                                                                                                                                     80.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         25pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                       0,
                                                                                                                                                                                                                                                                                                                                                                                                                                       Score 41; DB 22
Pred. No. 0.014;
                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 22; Length 17;
                                                                                                                                                                                                                                                                                                                                                                                                                       <u>ن</u>
                                                                                                                                 TSE; infection,
                                                                                                                                                                                                                                                                                                                                                                                                                       0:
                                                                                                                                                                                                                                                                                                                                                                                                                       Gaps
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RESULT 15 AAR38042

AAR38042 standard; protein; 22 AA.

14-OCT-1993 AAR38042;

(first entry)

Bovine prion protein region F #2.

Antigen; prion; protein; region; frame shift; repeat; mutation; PrPC; FSa; FSb; subfragment; antibody; treatment; spongiform encephalopathy; human; sheep; cattle; cellular binding; aggregation; mammal; scraple; immune system; PrPsc; ratio-inverso peptide; enzymatic degradation;

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Matches Query Match Best Local :

Local Similarity nes 8; Conserv

Conservative

80.4%;

Score 41; D Pred. No. 0. Mismatches

DB . 0.016; 23; Çī

Length 19; Indels

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Gaps

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CC thermostable proteolytic enzyme; (2) an antibody (II) specific for a CC prion dimer which does not bind to a prion monomer; and (3) a purified CC prion dimer. (M1) is useful for inactivating a TSE agent such as a prion. CC A TSE agent is Creutzfeldt-Jacob disease or its variant, kuru, fatal CC familial insomnia, Gerstmann-Straussler-Scheinker syndrome, bovine CC spongiform encephalopathy, scrapie, feline spongiform encephalopathy, (I) is CC useful for sterilising material contaminated with the TSE agent. A prion CC dimer is useful for examining a sample infected with or suspected to be CC infected by a prion protein, and for detecting prion infectivity, by detecting a prion dimer in the sample. A prion dimer, obtaining its contaminated with a prion dimer, obtaining its contained with a prion dimer, obtaining its contamined with a prion dimer, obtaining its contract which contains (II), and isolating (II) from the extract. The CC extract which contains (II), and isolating (II) from the preparation. (M1) CC and (I) are useful for inactivating TSE agents in potentially contaminated clinical waste and culled animal material. (M1) is useful CC for sterilising larger surface areas of apparatus, operating tables or CC even walls of rooms. The present sequence represents a prion mimetic CC peptide which is used in an example from the present invention in the preparation of antibodies including dimer preferential antibodies.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               08-JAN-2002; 2002WO-GB00052.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WO200253723-A2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  transmissible spongiform encephalopathy (TSE) agent comprising expothe TSE agent to a thermostable proteolytic enzyme. Also described:
(1) a composition (I) for inactivating a TSE agent, comprising a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The present invention describes a method (M1)
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26-FEB-2001; 2001GB-0004696.
Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         thermostable proteolytic enzyme
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      as Creutzfeldt-Jacob disease, scrapie, kuru or
Gerstmann-Straussler-Scheinker syndrome involves exposing agent to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Inactivating transmissible spongiform encephalopathy (TSE) agent such
as Creutzfeldt-Jacob disease, scrapie, kuru or
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Raven NDH
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resistance

i/Search completed: March 24, 2003, 17:19:41
 Job time : 37.1042 secs

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20
                                                                                                                                    Matches
                                                                                                Query Match
                                                                                                                                                           The sequences given in AAR38041-48 represent polypeptides which are derived from an antigenic site, region F, of a prion protein. Prion proteins comprise six regions of interest (A-F), and two related frame shift peptides sequences caused by a repeating section in region E having a nucleic acid coding sequence frame shift mutation of +1 (FSa) or -1 (FSb). These peptides (see also AAR38041-48) and antibodies raised against these may be used to treat or prevent spongiform encephalopathy in humans, sheep or cattle. They can be used to block cellular binding and aggregation of prion proteins and to stimulate the mammalian immune system. These peptides may be used to distinguish between the normal form of prion protein (PrPc) and the scrapic-associated form (PrPsc). These peptides may include rare or synthetic amino acids or a ratio-inverso peptide modification to improve resistance to enzymatic degradation.
                                                                                                                                  Sequence
                                                                                                                                                                                                                                                                                                                                                                                                                                                        New polypeptide(s) contg. antigenic site of prion protein - useful for treatment and diagnosis of mammalian encephalopathies e.g. Creutzfeld-Jacob disease and kuru
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M1sc-d1fference 21
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                                                                                                                                                                                                                                                                                                                                                                                                                          Claim 28; Page 74; 82pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       03-DEC-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Misc-difference
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  WPI; 1993-196994/24
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Fishleigh RV, Mee RP,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    03-DEC-1991;
10-JUL-1992;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         10-JUN-1993.
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                                                     80.4%;
Local Similarity 61.5%;
nes 8; Conservation
                              CXTQYXXESXAXY 13
                                                                                                                                 22 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    91GB-0025747.
92GB-0014663.
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                                                            Score 41; DB 14; Length 22;
Pred. No. 0.018;
0; Mismatches 5; Indels
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Maximum Match 100%
Listing first 45 summaries
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Maximum DB
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 Score
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seq length: 2000000000
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 protein search, using sw model
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1: pir1:*
2: pir2:*
3: pir3:*
4: pir4:*
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 CXTQYXXESXAXY 13
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UJH71H
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A53892
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S71041
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S7107048
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major prion PrP27-
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45	44	43	42	41	40	39	38	37	36	35	34	33	32	31	30
29	29	29	30	30	30	31	41	41	41	41	41	41	41	41	41
56.9	56.9	56.9	58.8	58.8	58.8	60.8	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4
639	392	392	758	499	339	938	264	264	260	257	256	256	256	254	254
N	N	N	N	N	N	N	N	N	N	N	N	N	N	N	Ν
S23569	T19867	151422	865169	D69735	JC7509	н83050	A54330	S37137	S53629	JQ1900	A54281	S37149	JU0268	A23544	B34759
gag polyprotein ho	hypothetical prote	glutamine syntheta	hypothetical prote	xylulose kinase xy	glycoprotein VI-1	probable ferredox1	major prion protei	prion protein - gr	major prion protei		major prion protei	prion protein - go	prion	major prion protei	prion protein - go

ALIGNMENTS

C:Species: Oryotolagus cuniculus (domestic rabbit) C:Species: Oryotolagus cuniculus (domestic rabbit) C:Accession: JC6175 R:Loftus, B: Rogers, M. Gene 184, 215-219, 1997 A;Ritle: Characterization of a prion protein (PrP) gene from rabbit; a species v A;Reference number: JC6175; MUID:9718365; PMID:9031631 A;Recession: JC6175 A;Rocession: JC6175 A;Rocession: JC6175 A;Rocession: JC6175 A;Rocession: JC6175 A;Rocession: JC6175 A;Rocession: JC6175 A;Cross-references: GB:U28334; NID:91490412; PIDN:AAC48697.1; PID:91490413 C;Comment: This protein is a cellular protein, it is involved in the neurodegene C;Genetics: A;Gene: PrP C:Superfamily: major prion protein C;Keywords: disulfide bond; prion RESULT 1 JC6175 Query Match Best Local 82.4%; Score 42; DB 2; Pred. No. 0.021; Length 252; rabbit; a species with a PID:g1490413 in the neurodegenerativ

Db 213 CITQYQQESQAAY 225
RESULT 2
ОЈНУІН
major prion PrP-Sc protein precursor - golden hamster
N; Contains: scrapie amyloid protein (PrP27-30)
C; Species: Mesocricetus auratus (golden hamster)
C;Date: 04-Dec-1986 #sequence_revision 12-Apr-1996 #text_change 28-Jan-2000
C;Accession: I48168; A03133; I48167; S02520; A36019; A40665
R; Basler, K.; Oesch, B.; Scott, M.G.; Westaway, D.; Waelchli, M.; Groth, D.F.; McKinl
Cell 46, 417-428, 1986
A; Title: Scrapie and cellular PrP isoforms are encoded by the same chromosomal gene.
A; Reference number: I48168; MUID: 86272089; PMID: 2873895
A; Accession: I48168
A; Molecule type: DNA
A; Residues: 1-254 <res></res>
A; Cross-references: GB: M14054; NID: g191425; PIDN: AAA37091.1; PID: g387076
R;Oesch, B.; Westaway, D.; Walchli, M.; McKinley, M.P.; Kent, S.B.H.; Aebersold, R.;
Cell 40, 735-746, 1985
A; Title: A cellular gene encodes scrapie PrP 27-30 protein.
A; Reference number: A03133; MUID: 85176927; PMID: 2859120
A;Accession: A03133
A; Molecule type: mRNA
A; Residues: 15-254 <oes></oes>
A;Cross-references: GB:K02234; NID:g191429; PIDN:AAA37092.1; PID:g191430

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1 CXTQYXXESXAXY 13

Matches

Local Similarity hes 8; Conserv

Conservative

0;

Mismatches

Indels

0

Gaps

0

prion

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A; Molecule type: protein
A; Residues: 23-24,'X',26-36,'X',38-55 <TUR>
A; Residues: 35-24, Padgett, M.P.; Ceroni, M.; Piccardo, P.; Zopf, D.; Gajdusek, D.C.
A; Title: Molecular mass, biochemical composition, and physicochemical behavior of the in
A; Reference number: A36019; MUID:90349618; PMID:1974720
A; Reference number: A36019; MUID:90349618; PMID:1974720
A; Residues: '5','24-32 <CSAF>
A; 
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                                                                                                                                                                                                                                                                                                                                                                   R;Robakis, N.K.; Sawh, P.R.; Wolfe, G.C.; Rubenstein, R.; Carp, R.I.; Innis, M.A. Proc. Natl. Acad. Sci. U.S.A. 83, 6377-6381, 1986
A;Title: Isolation of a cDNA clone encoding the leader peptide of prion protein and A;Reference number: A23545; MUID:86313584; PMID:3529083
A;Accession: A23545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    A:Status: preliminary; translated from GB/EMBL/DDBJ
A:Molecule type: mRNA
A:Molecule type: mRNA
A;Residues: 12-254 <RE2>
A;Cross-references: GB:M37381; NID:g191422; PIDN:AAA37090.1; PID:g191423
A;Cross-references: GB:M37381; NID:g191422; PIDN:AAA37090.1; PID:g191423
A;Turk, E.; Teplow, D.B.; Hood, L.E.; Prusiner, S.B.
Eur. J. Biochem. 176, 21-30, 1998
A;Title: Purification and properties of the cellular and scrapie hamster prion A;Reference number: S02519; MUID:88329062; PMID:3138115
A;Accession: S02520
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          major prion PrP27-30 protein - hamster
c;Species: Cricetinae gen. sp. (hamster)
C;Date: 29-Aug-1987 #sequence_revision 29-Aug-1987 #text_change 18-Jun-1993,
C;Accession: A23545
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 3
A23545
                                                                                                                                                                                                                                             C; Superfamily: major prion protein
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Int. Rev. Neurobiol. 28, 1-57, 1986
A;Title: Biology and structure of scrapie prions.
A;Reference number: I48167; MUID:87108309; PMID:3100471
A;Accession: I48167
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A; Contents: annotation; chemical (not conformational) identity of PrPSc and PrPC molecu A; Note: no chemical difference between cellular (PrPC) and scraple (PrPSc) forms of this
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           A; Note: this mRNA, isolated from the brain of scraple-infected hamster, is the product of R; Stahl, N.; Baldwin, M.A.; Teplow, D.B.; Hood, L.; Gibson, B.W.; Burlingame, A.L.; Prus Biochemistry 32, 1991-2002, 1993
                                                                                       Best Loc
Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         B2.4%;
Local Similarity 61.5%;
les 8; Conservation
                                                                                                                          Local
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CXTQYXXESXAXY 13
                                                                                   Similarity
8; Conserv
                                                                                       Conservative
                                                                                                                      82.4%;
61.5%;
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0; Mismatches
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A53892
A53892
prion-related protein - rat (fragment)
C;Species: Rattus norvegicus (Norway rat)
C;Species: Rattus norvegicus (Norway rat)
C;Date: 07-Oct-1994 #sequence_revision 07-Oct-1994 #text_change 13-Aug-1999
C;Accession: A53892
R;Liao, Y.C.; Tokes, Z; Lim, E; Lackey, A; Woo, C.H.; Button, J.D.; Clawson, Lab. Invest. 57, 370-374, 1987
A;Title: Cloning of rat "prion-related protein" cDNA.
A;Reference number: A53892; MUID:88037055; PMID:2889848
A;Accession: A53892
A;Accession: A53892
A;Status: preliminary
A;Molecule type: mRNA
A;Residues: 1-226 <LIA>
A;Cross-references: GB:M20313; NID:g206391; PIDN:AAA41947.1; PID:g206392
C;Superfamily: major prion protein
                 major prion protein - douroucoull (fragment)
C;Species: Actus trivirgatus (douroucoull, night monkey, owl monkey)
C;Date: 28-Oct-1996 #sequence_revision 07-Feb-1997 #text_change 13-Aug-1999
C;Accession: $53633; $71042
R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.
J. Mol. Biol. 245, 362-374, 1995
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C;Superfamily: major prion protein
C;Keywords: amyloid; brain; 9lycoprotein; lipoprotein; prion; scrapie; transmembrane
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submitted to the EMBL Data Library, April 1994
A:Reference number: S71041
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A;Status: nucleic acid sequence not shown
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   A;Cross-references: EMBL:U08309; NID:g474376; PIDN:AAC50097.1; PID:g474377 R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B. J. Mol. Biol. 245, 362-374, 1995
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A; Residues: 1-232 <SCH>
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A; Residues: 1-194, 'R', 196-231 <SCW>
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Best Local (
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Pred. No. 0.033;
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Pred. No. 0.
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PMID:7837269
primates
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A;Molecule type: DNA
A;Residues: 1-202, 'E',204-239 <SCW>
A;Residues: 1-202, 'E',204-239 <SCW>
A;Cross-references: EMBL:U08293; NID:g474344; PIDN:AAC50082.1; PID:g474345
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
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submitted to the EMBL Data Library, April 1994
A:Reference number: S71041
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A:Reference number: S53614; MUID:95139066; PMID:7837269
A:Accession: S53621
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C:Species: Papio sphinx, Mandrillus sphinx (mandrill)
C:Date: 27-Ct-1996 #sequence_revision 14-Feb-1997 #t
                                                                                         R:Schatzl, H.M.
submitted to the EMBL Data Library, April 1994
A:Reference number: S71041
A:Accession: S71048
                                                                                                                                                                                C;Species: Callicebus moloch
C;Date: 27-Oct-1996 #sequence_revision
C;Accession: S71048; S53632
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C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane pro
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A; Residues: 1-241 <SCH>
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A;Accession: S53633
A;Status: nucleic acid sequence not shown
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A; Residues: 1-203, 'R', 205-240 <SCW>
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A; Residues: 1-239 <SCH>
A;Cross-references: EMBL:U08312; NID:g475585; PIDN:AAC50100.1; PID:g475586 R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.
                                             A; Molecule type: DNA
A; Residues: 1-241 <SCH>
                                                                                                                                                                                                                                                      major prion protein - Callicebus moloch (fragment)
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nes 8; Conserv
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Similarity 61.5%;
8; Conservation
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Pred. No. 0.035;
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Pred. No. 0.034;
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R;Schatzl, H.M.

R;Accession: S71041

R;Accession: S71043

A;Accession: S71044

A;Accession: S71044
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A; Title: Prion protein gene variation among primates.
A; Reference number: $53614; MUID:95139066; PMID:7837269
A; Accession: $53632
A; Status: nucleic acid sequence not shown
A; Molecule type: DNA
A; Residues: 1-203, 'R', 205-240 < SCW>
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C;Date: 28-Oct-1996 #sequence_revision 07-Feb-1997 #text_change 13-Aug-1999
C;Accession: S53627; S71043
R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B.
J. Mol. Biol. 245, 362-374, 1995
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C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane
    A;Title: Prion protein gene variation among primates. A;Reference number: S53614; MUID:95139066; PMID:7837269 A;Accession: S53628 A;Status: nucleic acid sequence not shown
                                                                                                                                                             A;Cross-references: EMBL:U08292; NID:g474342; PIDN:AAC50081.1; PID:g474343 R;Schaetzl, H.M.; da Costa, M.; Taylor, L.; Cohen, F.E.; Prusiner, S.B. J. Mol. Biol. 245, 362-374, 1995
                                                                                                                                                                                                                                                                                                            A; Reference number: S71
A; Accession: S71045
A; Molecule type: DNA
A; Residues: 1-245 <SCH>
                                                                                                                                                                                                                                                                                                                                                                                                                                             submitted to the EMBL Data Library, A; Reference number: S71041
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              major prion protein - Cercopithecus dianae
C;Species: Cercopithecus dianae
C;Date: 14-Feb-1997 #sequence_revision 14-Feb-1997 #text_change 13-Aug-1999
C;Accession: S71045; S53628
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A; Residues: 1-245 <SCH>
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Best Local Similarity
Thes 8; Conserve
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Best Local (
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acid sequence not shown
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Pred. No. 0.035;
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Rivervenakova, L.; Brown, P.; Goldfarb, L.G.; Nagle, J.; Pettrone, K.; Rubenstein, R.; Proc. Natl. Acad. Sci. U.S.A. 91, 12159-12162, 1994.
A;Title: Infectious amyloid precursor gene sequences in primates used for experimental A;Reference number: I36907; MUID:95083661; PMID:7991600
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                                                                                                                                                 A;Molecule type: DNA
A;Residues: 1-209,'E',211-252 <SCW>
A;Residues: 1-209,'E',211-252 <SCW>
A;Cross-references: EMBL:008304; NID:g474366; PIDN:AAC50092.1; PID:g474367
C;Superfamily: major prion protein
C;Superfamily: major brion protein; lipoprotein; prion; scrapie; transmembrane pro
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 A;Cross-references: EMBL:U15165; NID:g595852; PIDN:AAA68636.1; PID:g595853 C;Superfamily: major prion protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       C;Species: Saimiri sciureus (C;Date: 31-May-1996 #sequence_revision C;Accession: 1848 Prown. P.; Goldfarb,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A;Cross-references: EMBL:U08292
C;Superfamily: major prion protein
C;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane
                                                                                                                                                                                                                                                                                                                                                                                                                                   R;Schaetzl, H.M.; db Costa, M.; Taylor, L.; Cohen, F.E.; J. Mol. Biol. 245, 362-374, 1995
A;Title: Prion protein gene variation among primates. A;Reference number: $53614; MUID:95139066; PMID:7837269
A;Accession: $53634
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C; Species: Saimiri sciureus (common squirrel monkey)
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A; Residues: 8-10, 'L', 12-202, 'R', 204-239 <SCW>
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A; Residues: 1-252 <RES>
                                                                                                                                                                                                                                                                                                                                                      A; Cross-references: EMBL: U08304
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Species: Callithrix jacchus (common marmoset)
Date: 28-Oct-1996 #sequence_revision 07-Feb-1997 #text_change 13-Aug-1999;Accession: S53634; S71047
                                                                                                                         Query Match
                                                                                                                                                                                                                                                                    Accession: S71047
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Residues: 1-252 <SCH>
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Best Local :
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                                        CXTQYXXESXAXY 13
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Pred. No.
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Pred. No. 0.035;
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Pred. No.
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A;Reference number: S53614;
A;Accession: S53631
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A; Residues: 1-80,89-253 < PUC>
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A; Residues: 1-253 < KRE>
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A; Residues: 1-252 <SCH>
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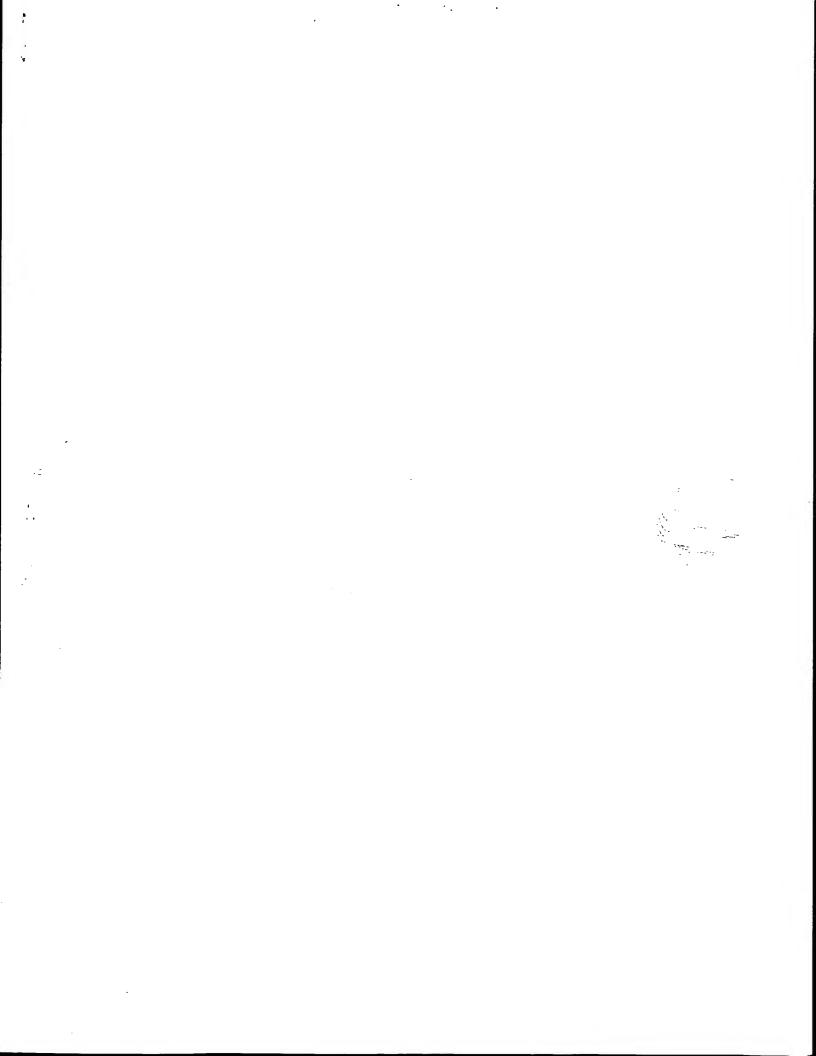
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A; Title: Amyloid protein of Gerstmann-Straeussler-Scheinker disease (Indiana kindred) A; Reference number: S14078; MUID:91160504; PMID:1672107
A; Accession: S14078
                                                                                                              A; Molecule type: mRNA
A; Residues: 8-117,119-253 <LIA>
A; Residues: 8-117,119-253 <LIA>
A; Cross-references: GB:D00015; NID:g220015; PIDN:BAA00011.1; PID:g220016; GB:M13667; R; Tagliavini, F.; Prelli, F.; Ghiso, J.; Bugiani, O.; Serban, D.; Prusiner, S.B.; FalEMBO J. 10, 513-519, 1991
                                                                                                                                                                                                                                                                                                              A;Note: the deletion may be a polymorphism; the alterna R;Liao, Y.C.J.; Lebo, R.V.; Clawson, G.A.; Smuckler, E. Science 233, 364-367, 1986
A;Reference number: A05017; MUID:86261778; PMID:3014653
A;Accession: A05017
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C;Date: 25-Oct-1987 #sequence_revision 12-Apr-1996 #text_change 16-Jun-2000
C;Accession: A24173; A40372; A05017; S14078; I54322; I68597; I58135; I59184; I79633;
R;Kretzschmar, H.A.; Stowring, L.E.; Westaway, D.; Stubblebine, W.H.; Prusiner, S.B.
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A;Molecule type: DNA
A;Residues: 17209, 'E',211-252 <SCW>
C;Superfamily: major prion protein prion; prion; scrapie; transmembrane c;Keywords: amyloid; brain; glycoprotein; lipoprotein; prion; scrapie; transmembrane
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A;Title: Genomic structure of the human prion protein gene. A;Reference number: A40372; MUID:91328137; PMID:1678248
A;Accession: A40372
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A;Title: Molecular cloning of a human prion protein cDNA.
A;Reference number: A24173; MUID:86300093; PMID:3755672
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A;Residues: 1-80,89-200 A;Residues: 1-80,89-200 A;Cross-references: GB:X83416; NID:g747846; PIDN:CAA58442.1; PID:g747847
A;Cross-references: GB:X83416; NID:g747846; PIDN:CAA58442.1; PID:g747847
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N;Alternate names: 11K amyloid protein; 27-30K sialoglycoprotein; PrP 27-30;
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C;Date: 28-Oct_1996 #sequence_revision 07-Feb-1997 #text_change
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A; Status: not compared with conceptual translation
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$53614; MUID:95139066; PMID:7837269
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A; Introns: #status abbent
A; Note: one intron occurs before the initiator codon
A; Note: one intron occurs before the initiator codon
A; Note: this gene is associated with Creutzfeld-Jakob disease (CJD), Gerstmann-Strausler
C; Superfamily: major prion protein
C; Keywords: amyloid; blocked carboxyl end; brain; glycoprotein; lipoprotein; phosphatidy
E; 1-22/Domain: signal sequence #status predicted <SIG>
F; 23-230/Product: major prion protein #status predicted <MAT>
F; 54-92/Region: 8-residue repeats (P-H-G-G-G-W-G-Q)
F; 112-134/Domain: transmembrane #status predicted <TM1>
F; 231-253/Domain: carboxyl-terminal propeptide #status predicted <CTP>
F; 179-214/Disulfide bonds: #status predicted
F; 181,197/Binding site: carbohydrate (Asn) (covalent) #status predicted
F; 181,197/Binding site: GPI-anchor ethanolamine amidated carboxyl end (Ser) (in mature form
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A;Cross-references: GB:S80539; NID:g244698; PIDN:AAB21334.1; PID:g244699
R;Goldfarb, L.G.; Brown, P.; McCombie, W.R.; Goldgaber, D.; Swergold, G.D.; Wills, P.R.
Proc. Natl. Acad. Sci. U.S.A. 88, 10926-10930, 1991
A;Title: Transmissible familial Creutzfeldt-Jakob disease associated with five, seven, A;Reference number: I59184; MUID:92073400; PMID:1683708
A;Accession: I59184
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A; Molecule type: protein
A; Residues: 58-72, 'X', 74-76, 'XXX', 79, 'XXX', 83-86; 111-128, 'V', 130-150 <TAG>
R; Diedrich, J.F.; Knopman, D.S.; List, J.F.; Olson, K.; Frey, W.H.
Hum. Mol. Genet. 1, 443-444, 1992
A; Title: Deletion in the prion protein gene in a demented patient.
A; Reference number: I54322; MUID:93250789; PMID:1363802
A; Accession: I54322
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R;Brown, P.; Goldfarb, L.G.; McCombie, W.R.; Nieto, A.; Squillacote, D.; Sheremata, W.; Neurology 42, 422-427, 1992
A;Title: Atypical Creutzfeldt-Jakob disease in an American family with an insert mutatic A;Reference number: I58135; MUID:92140671; PMID:1736177
A;Accession: I58135
                                                                                                                                                                                                                         major prion protein precursor - gorilla (c;Specles: Gorilla gorilla (gorilla) (c;Specles: Gorilla gorilla (gorilla) (c;Date: 31-May-1996 *sequence_revision 31-May-1996 *text_change 13-Aug-1999 (c;Accession: I37032 (c;Ac
A;Title: Infectious amyloid precursor gene sequences in primates used for experimental A;Reference number: I36907; MUID:95083661; PMID:7991600
A;Accession: I37032
A;Status: preliminary; translated from GB/EMBL/DDBJ
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A; Residues: 8-240 <R
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A;Accession: I68597
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A; Residues: 9-83,92-240 <RES>
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Residues: 60-67 <GOL>
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A;Molecule type: DNA
A;Residues: 1-253 <RES>
A;Cross-references: EMBL:U15166; NID:g563208; PIDN:AAA68633.1; PID:g563209
C;Superfamily: major prion protein
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214 CITQYERESQAYY 226
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Search completed: March 24, 2003, Job time: 17.25 secs

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95211 oryctolagus
P04273 mesocricetu
P40246 atteles geof
95145 cercocebus
95270 theropithec
P40248 callicebus
P40255 mandrillus
P40256 cercopithec
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P40257 cercocebus
P51446 atteles pani
P40247 callithrix
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P40252 gorilla gor
P04155 pong origina gor
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P40258 pan troglod
P40259 pan troglod
P40250 pongo pygma
P40257 presbytis f
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P604025 mus musculu
P13852 rattus norv
Q92013 sigmodon hi
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P47852 doccolleus
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AA TANDEM REPEATS OF P-H-G-G-G-W-G-

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                                                                       MEDLINE-91104789; PubMed-1980209; Stahl N., Baldwin M.A., Burlingame A.L., Prusiner S.B.; Stahl N., Baldwin M.A., Burlingame A.L., Prusiner S.B.; "Identification of glycoinositol phospholipid linked and truncated forms of the scraple prion protein."; Blochemistry 29:8879-8884(1990).
Rogers M., Taraboulos A., Scott M., Groth D., Prusiner S.B.; "Intracellular accumulation of the cellular prion protein after mutagenesis of its Asn-linked glycosylation sites.";
                                                                                                                                                          MEDLINE-88027007; PubMed-2444340;
Stahl N., Borchelt D.R., Hasiao K
"Scrapit prion protein contains a
Cell 51:229-240(1987).
                                                                                                                                                                                                                                                        Turk E., Teplow D.B., Hood L.E., Pruse Florification and properties of the
                                     MEDLINE-92369621; PubMed-1983782;
                                                                                                                                       GPI - ANCHOR
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                                                                                                                                                                                                                                                prion
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                                                                                                                                                                                                                                                                                            CARBOHYDRATE-LINKAGE SITES,
                                                                                                                                                                                                                                                                                                                                              Basler K., Oesch B., Scott M., Westaway D., Waelchli M., Groth D.F., McKinley M.P., Prusiner S.B., Weissmann C.; "Scraple and cellular PrP isoforms are encoded by the same chromosomal gene."; cell 46:417-428(1986).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     REPEAT
REPEAT
                                               MUTAGENESIS OF CARBOHYDRATE-LINKAGE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Mesocricetus auratus (Golden hamster).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Cricetinae;
                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE OF 15-254 FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID-10036;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MEDLINE-86272089; PubMed-2873895;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                   cellular gene encodes scrapie PrP 27-30 protein.";
11 40:735-746(1985).
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8; Conser
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(Rel. 37, Last annotation update)
protein precursor (PrP) (PrP27-30) (PrP33-35C).
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83
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27432 MW;
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Pred. No.
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                                                                                                                                                                  K., Prusiner S.B.;
a phosphatidylinositol glycolipid.";
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0.0067;
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Hood L.E.,
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Query Match
Best Local S
Matches
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DISULFID
                                                        MUTAGEN
MUTAGEN
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REPEAT
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REPEAT
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                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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                                                                                                                                               DOMAIN
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                                                                                                                                                                                                                                      SIGNAL
                                                                                                                                                                                                                                                                             Pfam; PF00377; prion; PRINTS; PR00341; PRIO; SMART; SM00157; PRP;
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InterPro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MEDLINE-97439821; PubMed-9294167;
James T.L., Liu H., Ulyanov N.B., Farr-Jones S.,
Donne D.G., Kaneko K., Groth D., Mehlhorn I., Pru
Cohen F.E.;
                                                                                                                                                                                                     LIPID
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[7]
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           "Solution structure of a 142-residue recombinant prion protein corresponding to the infectious fragment of the scrapie isoforn Proc. Natl. Acad. Sci. U.S.A. 94:10086-10091(1997).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STRUCTURE BY NMR OF 90-231
                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIM INFECTED WITH DEGENERATIVE NEUROLOGICAL DISEASES SUCH AS KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                Brain;
   Similarity
8; Conser
                                              254 AA;
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60
68
76
84
183
   Conservative
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67
75
83
91
                                             27919
            82.4%;
61.5%;
                                              MW.
  0,
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            Score 42; I
Pred. No. 0
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                                                                                                                                                                             N-LINKED
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(GLCNAC. . .).
(GLCNAC. . .).
            DB 1;
0.0067
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Prusiner
                    Length 254;
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                                             CRC64;
  Indels
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S.B.,
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Gaps
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1 CXTQYXXESXAXY

RESULT 3
PRIO_ATEGE

01-FEB-1995 01-OCT-1996 Major prion

(Rel. 31, Created)
(Rel. 31, Last sequence update)
(Rel. 34, Last annotation update)
protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).

01-FEB-1995 P40246; PRIO_ATEGE

STANDARD;

PRT;

232

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MEDILINE-95139066; PubMed-7837269;
Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;
"Prion protein gene variation among primates.";
J. Mol. Biol. 245:362-374(1995).
-1- EUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-1- SUBBUNIT: PRP HAS A TENDENCY TO AGGREGATE VIELDING POLYMERS CALLED

SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE QUANTITY IN THE BRAIN OF HUMANS AND CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.

"RODS"

SIMILARITY: BELONGS TO THE PRION FAMILY.

Ateles geoffroyi (Black-handed spider monkey). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Mammalia; Eutheria; Primates; Platyrrhini; Cebidae;

Euteleostomi; Atelinae; Ateles

NCBI_TaxID-9509;

SEQUENCE FROM N.A.

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214 CTTQYQKESQAYY 226

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                  Matches
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CARBOHYD
CARBOHYD
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                                                    SEQUENCE
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                                                                                                                                                                              SIGNAL
                                                                                                                                                                                                        PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
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SMART; SM00157; PRP; 1.
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HSSP; P04156; LEIG.
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                           Local
 1 CXTQYXXESXAXY 13
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                           Similarity
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232 AA;
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215
214
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163
165
181
                  Conservative
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58
68
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67
75
                         80.4%;
                                                     25596 MW;
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                 0,
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N-LINKED
4 X 8 AA
                 Score 41; DB 1;
Pred. No. 0.011;
0; Mismatches
                                                                       <u>*</u> 3210
                                                                                                                                         MAJOR PRION PROTEIN.
REMOVED IN MATURE FORM (BY SIMILARITY).
GPI-ANCHOR (BY SIMILARITY).
BY SIMILARITY.
                                                     0E2D75F04C05CC4A CRC64;
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                                                                                                                  AA TANDEM REPEATS OF P-H-G-G-G-W-G-
                                                                                                                          (GLCNAC. . .) (POTENTIAL).
                                   Length 232;
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                 Gaps
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1 CXTQYXXESXAXY 13

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01-NOV-1997 (Rel. 35, Created)
01-NOV-1997 (Rel. 35, Last sequence update)
15-JUN-2002 (Rel. 41, Last annotation update)
15-JUN-2002 (Rel. 41, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                der Kuyl A.C., Dekker J.T., Goudsmit J.;

Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.

SUBMITTED THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.

-I- SUBBULIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                        LIPID
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                                                                                           REPEAT
REPEAT
                                                                                                                                                                                                                                                     CHAIN
                                                                                                                                                                                                                                                                                                                    PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                                                                                                                                                                                                                                                                                                   Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                  EMBL; U75382; AAB50629.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Macaca sylvanus (Barbary ape).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cercocebus aterrimus, and
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NCBI_TaxID=36222, 9546;
                                                               SEQUENCE
                                                                                                                                                                          CARBOHYD
                                                                                                                                                                                         CARBOHYD
                                                                                                                                                                                                                                                                       SIGNAL
                                                                                                                                                                                                                                                                                    NON_TER
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                                                                                                                            REPEAT
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                                                                                                                                                                                                                                                                                                       Prion;
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                Local
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                                                                                                                                                                                                                                                                                                       Brain;
                Similarity
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                                                               238
   Conservative
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                                                                                            52
68
                                                             26123 MW;
              80.4%;
                                                                                                                                                                                                                                                                                                     GPI-anchor; Repeat;
Score 41; DB 1;
Pred. No. 0.011;
0; Mismatches
                                                                                                                                       N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
4 X 8 AA TANDEM REPEATS OF P-H-G-G-C.
                                                                                                                                                                                                      REMOVED IN MATURE FORM (BY SIMILARITY). GPI-ANCHOR (BY SIMILARITY). Y
BY SIMILARITY.
                                                                                                                                                                                                                                                       MAJOR PRION PROTEIN
                                                                                                                                                                                                                                                                       BY SIMILARITY.
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                                                               5F59A3EBC3E3531B CRC64;
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                               Length 238;
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELLT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                              SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 EMBL; U75383; AAB50630.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          der Kuyl A.C., Dekker J.T., Goudsmit J.;
Submitted (NOV-1996) to the EMBL/GenBank/DDBJ databases.
-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
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01-NOV-1997 (Rel. 35, Last sequence update)
01-NOV-1997 (Rel. 35, Last sequence update)
Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Prion; Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR000817; Prion. Pfam; PF00377; prion; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRIO_THEGE Q95270;
199 CITQYQKESQAYY 213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     -1- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS
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                                                                                                                                        Local Similarity
                                                  1 CXTQYXXESXAXY 13
                                                                                                             Conservative
                                                                                                                                                                                                                                                                                       53
64
69
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>238
199
166
182
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                                                                                                                                  80.4%;
                                                                                                                                                                                                                              26104 MW;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          GPI-anchor; Repeat;
                                                                                                                                  Score 41; DB 1; Length 238; Pred. No. 0.011;
                                                                                                                                                                                                                                                                                                                                                                                                                MAJOR PRION PROTEIN.
BY SIMILARITY.
N-LINKED (GLCNAC. ..) (POTENTIAL).
N-LINKED (GLCNAC. ..) (POTENTIAL).
A X 8 AA TANDEM REPEATS OF P-H-G-G-W-G-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     BY SIMILARITY.
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                                                                                                             Mismatches
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206 CITQYEKESQAYY 218

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ID PRIO_A
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AC PAIN
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                                                                                  Matches
                                                                                                                                                            Query Match
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"Prion protein gene variation among primates.";
J. Mol. Biol. 245:362-374(1995).

-i- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.

-I- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
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Pfam; PF00377; prion; 1.
SMART; SM00157; PRP; 1.
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                                                                                                                                                                                                                                          SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
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01-FEB-1995 (Rel. 31, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CARBOHYD
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Aotus trivirgatus (Night monkey) (Douroucouli).
Eukaryota; Metazoa; Chordata; Craniata; Vertebr
Mammalia; Eutheria; Primates; Platyrrhini; Cebi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Major prion protein precursor (PrP) (PrP27-30)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Prion; Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NCBI_TaxID=9505;
1 CXTQYXXESXAXY 13
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                                                                                                                   Similarity
                                                                                                                                                                                                                                          239 AA;
                                                                              Conservative
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50
58
76
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                                                                                                                                                                                                                                      26246 MW;
                                                                                                                       80.4%;
                                                                                                                   . 5e;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MAJOR PRION PROTEIN.
BY SIMILARITY.
N-LINKED (GLCNAC. . .) (FOR THE PROPERTY OF THE PROPERTY O
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                                                                                                      Score 41; DB 1;
Pred. No. 0.011;
                                                                                                                                                                                                                                  2EFB77E354B7024A CRC64;
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                                                                         Mismatches
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                                                                                                                                                   Length 239;
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Best Local
                                                                                                        Matches
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01-FEB-1995
01-FEB-1995
01-NOV-1995
                                                                                                                                                                                                                                                                                                                                                                                                                                                 CARBOHYD
CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          MEDLINE=95139066; PubMed=7837269;
Schatzl H.M., Dacosta M., Taylor L., Cohen F.E.
"Prion protein gene variation among primates.";
J. Mol. Biol. 245:362-374(1995).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Callicebus moloch (Dusky titi).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Primates; Platyrrhini; Cebidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Pfam; PF00377; prion; SMART; SM00157; PRP; 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; U08312; AAC50100.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
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207
                                                 1 CXTQYXXESXAXY 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DECEMERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASES (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
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FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS. SUBURIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
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CITOYEKESQAYY 219
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Brain;
                                                                                                   8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              IPR000817; Prion.
                                                                                                                                                                                                            241 AA;
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(Rel. 31, Last sequence update)
(Rel. 32, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                protein precursor (PrP) (PrP27-30)
                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    STANDARD;
                                                                                                                                                                                                                                                                                                                                                                                                                        84
                                                                                                                                                                                                            26373 MW;
                                                                                                                               80.4%;
                                                                                                                                                                                                                                                           N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
5 X 8 AA TANDEM REPEATS OF P-H-G-G-W-G-Q.
1.
2.
3.
4.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            GPI-anchor; Repeat; Signal.
                                                                                                                               Pred
                                                                                                                                                  Score 41;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MAJOR PRION PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
                                                                                                                                                                                                            C6D2013EE7CAEC93 CRC64;
                                                                                                      Mismatches
                                                                                                                               . No. 0.011;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Cohen F.E., Prusiner S.B.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    241 AA
                                                                                                                                                       DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PrP33-35C) (Fragment).
                                                                                                                                                       Length 241;
                                                                                                   Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Euteleostomí;
Callicebinae;
                                                                                                   0;
                                                                                                   Gaps
                                                                                                   0
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RESULT 8
PRIO_MANSP
 Š
                                Query Match
Best Local S
Matches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-FEB-1995
01-FEB-1995
01-OCT-1996
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;
"Prion protein gene variation among primates.";
J. Mol. Biol. 245:362-374(1995).
-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                              NON_TER
SEQUENCE
                                                                                                                                                                                                                                                                                                                        CHAIN
                                                                                                                                               REPEAT
REPEAT
                                                                                                                                                                            REPEAT
REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics in the European Bioinformatics in There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Cercopithecinae; NCBI_TaxID=9561;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Eutele
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
Cercopithecinae; Mandrillus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Mandrillus sphinx (Mandrill) (Papio sphinx).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PRNP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Major prion
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P40255;
                                                                                                                                                                                                                                                                                                                                                                         Prion;
                                                                                                                                                                                                                                                                                                                                                                                                                         InterPro; IPR000817; Pr
Pfam; PF00377; Prion; 1
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MEDLINE=95139066; PubMed=7837269;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              PRIO_MANSP
                                                                                                                                                                                                                             DOMAIN
                                                                                                                                                                                                                                              CARBOHYD
                                                                                                                                                                                                                                                               CARBOHYD
                                                                                                                                                                                                                                                                            DISULFID
                                                                                                                                                                                                                                                                                                                                             SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                         PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        EMBL; U08303; AAC50091.1; -. HSSP; P04925; 1AG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       or send an email to license@isb-sib.ch).
                                                                                                                              REPEAT
                                                                                                                                                                                                                                                                                             LIPID
                                                                                                                                                                                                                                                                                                             PROPEP
                                                                                                                                                                                                                                                                                                                                                            NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AL ANYMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES (CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TRE), ETC.
CXTQYXXESXAXY
                                                                                                                                                                                                                                                                                                                                                                           Brain;
                                Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                           IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Rel. 31, Created)
(Rel. 31, Last sequence update)
(Rel. 34, Last annotation update)
protein precursor (PrP) (PrP27-30)
                                 Conservative
                                                                                                A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                STANDARD;
                                                                                                                                                                                                                                                                                                                                                                         lycoprotein;
   13
                                                                                                                                                                                                                           190
84
                                                                                                                                52
68
76
84
                                                                                                26398 MW;
                                               80.4%;
                                0;
                                                                                                                                                                                                                                                                                                                                                                         GPI-anchor; Repeat; Signal
                                               Score 41; I
Pred. No. 0.
                                                                                                                                                                                                                          N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
5 X 8 AA TANDEM REPEATS OF P-H-G-G-W-G-
                                                                                                                                                                                                                                                                                        REMOVED IN MATURE FORM (BY SIMILARITY).
GPI-ANCHOR (BY SIMILARITY).
                                                                                                                                                                                                                                                                            BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                             BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                            MAJOR PRION PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                                                                                E539D84E2E2B59DE CRC64;
                                Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                241 AA
                                               DB 1;
0.011;
                                                            Length 241;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (PrP33-35C) (Fragment).
                                Indels
                                0,
                                Gaps
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207 CITQYEKESQAYY

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RESULT 9
PRIO_CERAS
IDIO_CAC
PA0250
DT 01-FEB
DT 01-FEB
DT 01-FEB
DT 01-FEB
DT 01-FEB
OC Cercop
OC Eukary
OC CACCO
CCCCO
RLI J. Mammal
RR MEDLII
RR EMBI
DR EMBI
DR FFAN
CC -1- F
CC -1-
S
                                                                Matches
                                                                                        Query Match
Best Local :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.

MEDLINE-95139066; PubMed-7837269;

Schatzl H.M., Dacosta M., Taylor L., Cohen F.E.,

"Prion protein gene variation among primates.";

J. Mol. Biol. 245:362-374(1995)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00377; prion; 1
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                EMBL; U08291; AAC50080.1; -. EMBL; U08292; AAC50081.1; -. HSSP; P04925; 1AG2. Interpro; IPR000817; Prion.
                                                                                                                                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cercopithecus aethiops (Green monkey) (Grivet), and Cercopithecus diana (Diana monkey).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                    CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                   DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PROPEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Cercopithecinae; Cercopithecus.
NCBI_TaxID=9534, 36224;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
16-OCT-2001 (Rel. 40, Last annotation update)
Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRIO_CERAE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      P40250
                                                                                     Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURD, CREUTZFELLDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC. SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               MOL. BIO1. 245:362-374(1995).

FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORWAL AND INFECTED CELLS.

SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brain;
                                                          Similarity
8; Conserv
                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STANDARD;
                                                                                  80.4%;
                                                                                                                                                                              26885
                                                                                                                                                                              X
E
                                                       Score 41; DB
Pred. No. 0.01
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       GPI-anchor; Repeat; Signal.
                                                                                                                                                                                                                                                                                                                                                 REMOVED IN MATURE FORM (BY SIMILARITY).

GPI-ANCHOR (BY SIMILARITY).

BY SIMILARITY.

N-LINKED (GLCNAC. . .) (POTENTIAL).

N-LINKED (GLCNAC. . .) (POTENTIAL).

4 X 8 AA TANDEM REPEATS OF P-H-G-G-G-W-G-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MAJOR PRION PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRT;
                                                                                                                                                                           D582B58E2726C99A CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               245
                                                                                  DB 1;
0.011;
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                                                          Ģ
                                                                                                              Length 245;
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                                                    0;
                                                    Gaps
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PRIO_
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Query Match
Best Local S
Matches 8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          PRIO_CERMO STANDAKU;
PRIO_CERMO STANDAKU;

095172; 095173;
01-NCV-1997 (Rel. 35, Created)
01-NCV-1997 (Rel. 35, Last sequence update)
01-NCV-1998 (Rel. 36, Last annotation update)
15-JUL-1998 (Rel. 36, Last annotation update)
15-JUL-1998 (Rel. 36, Last annotation update)
                                                                                                            REPEAT
REPEAT
REPEAT
                                                                                                                                                                                                         LIPID
DISULFID
CARBOHYD
                                                              SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 der Kuyl A.C., Dekker J.T., Goudsmit J.;
Submitted (NOV-1996) to the EMEL/GenBank/DDBJ databases.
-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBBULT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
"RODS".
                                                                                                                                                                                                                                                                      CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
                                                                                REPEAT
                                                                                                                                                                                           CARBOHYD
                                                                                                                                                                                                                                                                                                                                 PROSITE; PS00291; PRION_1; PROSITE; PS00706; PRION_2;
                                                                                                                                                                                                                                                                                                                                                                                                              EMBL; U75386; AAB50625.1;
EMBL; U75387; AAB50626.1;
HSSP; P04925; IAG2.
                                                                                               REPEAT
                                                                                                                                                                           DOMAIN
                                                                                                                                                                                                                                                        PROPEP
                                                                                                                                                                                                                                                                                      SIGNAL
                                                                                                                                                                                                                                                                                                                       Prion;
                                                                                                                                                                                                                                                                                                                                                                  Pfam; PF00377; prion; SMART; SM00157; PRP;
                                                                                                                                                                                                                                                                                                                                                                                                 InterPro; IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                           or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cercopithecinae; Cercopithecus. NCBI_TaxID=36226, 36227;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Cercopithecus mona, and Cercopithecus neglectus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CERMO
                                                                                                                                                                                                                                                                                                       NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         NCBI_TaxID=36226,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 206 CITQYEKESQAYY 218
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRE IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREDTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                    Brain;
                Similarity
                                                              246
                                                                                                                                                                                                                                                                                                                    Glycoprotein;
                                                                                                                                                                         207
174
190
84
                                                            26900
                                                                                            52
68
76
              80.4%;
                                                              MW;
                                                                                                                                                                                                                                                                                                                  GPI-anchor; Repeat;
              Score 41; DB 1;
Pred. No. 0.011;
                                                                                                                                                                        N-LINKED
5 X 8 AA
                                                                                                                                                                                     BY SIMILARITY.
N-LINKED (GLCNAC. .) (POTENTIAL).
N-LINKED (GLCNAC. .) (POTENTIAL).
                                                                                                                                                                                                                                   MAJOR PRION PROTEIN.
REMOVED IN MATURE FORM (BY SIMILARITY).
GPI-ANCHOR (BY SIMILARITY).
                                                                                                                                                                                                                                                                                    BY SIMILARITY.
                                                              835D147CA2B4FDD3 CRC64;
                                                                                                                                                                         AA TANDEM REPEATS
                            Length 246;
                                                                                                                                                                                                                                                                                                                    Signal
                                                                                                                                                                         OF P-H-G-G-G-W-G-
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Conservative

0,

Mismatches

0;

Gaps

CXTQYXXESXAXY 13

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무
ş
              Query Match
Best Local Similarity
Thes 8; Conserv
                                                                                                                                                                                                     CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                           This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics in the European Bioinformatics in There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   01-NOV-1997
01-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Q95174;
01-NOV-1997
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CERPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            der Kuyl A.C., Dekker J.T., Goudsmit J.;
Submitted (NOV-1996) to the EMBLYDBJ databases.
-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE
HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIO_CERPA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Cercopithecinae; Cercopithecus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Cercopithecidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Cercopithecus patas.
                                                                                     SEQUENCE
                                                                                                                               REPEAT
                                                                                                                                                           REPEAT
                                                                                                                                                                                      DOMAIN
                                                                                                                                                                                                                                                                         CHAIN
                                                                                                                                                                                                                                                                                        SIGNAL
                                                                                                                                                                                                                                                                                                                                  PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                             Pfam; PF00377; prion; SMART; SM00157; PRP;
                                                                                                                                                                                                                                                                                                                                                                                                     EMBL; U75388; AAB50627.1;
HSSP; P04925; IAG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                 or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID-27677;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Major prion
                                                                                                                 REPEAT
                                                                                                                                            REPEAT
                                                                                                                                                                                                                                DISULFID
                                                                                                                                                                                                                                                             PROPEP
                                                                                                                                                                                                                                                                                                        NON_TER
                                                                                                                                                                                                                                                                                                                      Prion;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.
                                                                                                                                                                                                                                               TIPID
                                                                                                                                                                                                                                                                                                                                                                                           InterPro; IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  207
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURD CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONCIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITOYEKESQAYY 219
                                                                                                                                                                                                                                                                                                                      Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (Rel. 35, Last annotation update)
protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 (Rel. 35, Last sequence update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (Rel. 35, Created)
                             Conservative
                                                                                                                                                                                                                                                                                                                   Glycoprotein;
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                                                                                      26886 MW;
                                        80.4%;
61.5%;
                                                                                                                                                                                                                                                                                                                   GPI-anchor; Repeat; Signal.
                             0
                                         Score 41; I
Pred. No. 0.
                                                                                                                                                                                                                                         MAJOR PRION PROTEIN.
REMOVED IN MATURE FORM (BY SIMILARITY).
GPI-ANCHOR (BY SIMILARITY).
                                                                                                                                                                                                                   N-LINKED
                                                                                                                                                                                                                                BY SIMILARITY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             PRT;
                                                                                     D35D105BBEC53108 CRC64;
                                                                                                                                                                                                                                                                                         SIMILARITY.
                            Mismatches
                                                                                                                                                                                      AA TANDEM REPEATS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            246
                                                                                                                                                                                                     (GLCNAC. . .)
                                                        DB 1;
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                                           .011;
                                                       Length 246;
                             Indels
                                                                                                                                                                                                     (POTENTIAL).
                                                                                                                                                                                      OF P-H-G-G-G-W-G-
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PRIO_CERTOC
AC PRIO_CA
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Best Local
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REPEAT
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REPEAT
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBI outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE: PS00291; PRION_1;
PROSITE: PS00706; PRION_2;
Prion; Brain; Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Pfam; PF00377; prion; 1
SMART; SM00157; PRP; 1.
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                                                                                                                                                                      SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CHAIN
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                                                                                                                                                                                                                                                                                                                                                                                            CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                        DISULFID
                                                                                                                                                                                                                                                                                                                                             DOMAIN
                                                                                                                                                                                                                                                                                                                                                                       CARBOHYD
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1 CXTQYXXESXAXY 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-VAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CERTO
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                                                                  8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                non-profit
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(Rel. 35, Last sequence update)
(Rel. 35, Last annotation update)
protein precursor (PrP) (PrP27-30) (PrP33-35C) (Fragment).
                                                                                                                                                                      246
                                                                       Conservative
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                                                                                            80.4%;
                                                                                                                                                                      26914 MW;
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                                                                                                                                                                                                                          N-LINKED
5 x 8 AA
Q.
1.
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3.
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Pred. No. 0.
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GPI-ANCHOR (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             MAJOR PRION PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRT;
                                                                                                                                                                      F58679CBBEC5ADC7 CRC64;
                                                                          Mismatches
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                                                                                                                                                                                                                                                                                                                                             (GLCNAC. . .) (POTENTIAL). (GLCNAC. . .) (POTENTIAL). TANDEM REPEATS OF P-H-G-G-
                                                                                                                      DB 1;
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                                                                                               .011;
                                                                                                                      Length 246;
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OF P-H-G-G-G-W-G-
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CITQYEKESQAYY 219

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                                                           Query Match
     Matches
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REPEAT
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REPEAT
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"Infectious amyloid precursor gene sequences in primates used for experimental transmission of human spongiform encephalopathy.";
Proc. Natl. Acad. Sci. U.S.A. 91:12159-12162(1994).
-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.
-I- SUBUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                 SEQUENCE
                                                                                                                                                REPEAT
                                                                                                                                                                                                                                                                                                                                                                     CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CHAIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SIGNAL
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
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-i- DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZPELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                               DOMAIN
                                                                                                                                                                                                                                                                                                                                        CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   EMBL; U15164; AAA68634.1; -. HSSP; P04156; 1E1G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                    PROPEP
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Pfam; PF00377; prion; 1
PRINTS; PR00341; PRION.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 -1- SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ateles paniscus (Black spider monkey).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Platyrrhini; Cebidae; Atelinae; Ateles.
                                                                                                                                                                                                                                                                                                                                                                                              DISULFID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   InterPro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MEDLINE=95083661; PubMed=7991600
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRNP
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            PRIO_ATEPA
P51446;
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     Local Similarity
nes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             EQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         (Rel. 34,
(Rel. 34,
(Rel. 34,
     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      protein
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59
67
75
83
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STANDARD;
                                                                                                                                                                                                                                                                                                         229
229
229
229
213
180
196
                                                                                                        58
66
74
82
90
27718 MW;

    Last sequence update)
    Last annotation update)
    precursor (PrP) (PrP27-30) (PrP33-35C).

                            80.4%;
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BY SIMILARITY.
                                                                                                                                                                                                                                                                       BY SIMILARITY.

N'LINKED (GLCNAC. . .) (POTENTIAL).

N'LINKED (GLCNAC. . .) (POTENTIAL).

N'LINKED (GLCNAC. . .) (POTENTIAL).
                            Score 41; D
Pred. No. 0.
                                                                                                                                                                                                                                                                                                                                                                                                       MAJOR PRION PROTEIN.

REMOVED IN MATURE FORM (BY SIMILARITY).

GPI-ANCHOR (BY SIMILARITY).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  PRT;
                                                                                                        20EA38A42DCC56D1 CRC64;
     Mismatches
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                            .012;
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Indels
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 Query Match
Best Local Similarity
                                          REPEAT
SEQUENCE
                                                                     REPEAT
REPEAT
REPEAT
REPEAT
                                                                                                                                                         CARBOHYD
CARBOHYD
                                                                                                                                                                                                                                                                      PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       EMBL; U08304; AAC50092.1; -. HSSP; P04925; 1AG2. InterPro; IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                      This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way modified and this statement is not removed. Usage by and for commercial entities requires a license agreement (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A.
MEDLINE-95139066; PubMed-7837269;
Schatzl H.M., Dacosta M., Taylor L.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   P40247;
01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C).
                                                                                                                                            DOMAIN
                                                                                                                                                                                     DISULFID
                                                                                                                                                                                                                PROPEP
                                                                                                                                                                                                                              CHAIN
                                                                                                                                                                                                                                            SIGNAL
                                                                                                                                                                                                                                                                                                PRINTS; PR00341; PRION. SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                          InterPro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
                                                                                                                                                                                                                                                                                                                                                                                            or send an email to license@isb-sib.ch).
                                                                                                                                                                                                   LIPID
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Callithrix jacchus (Common marmoset).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NCBI_TaxID=9483;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Mammalia; Eutheria;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Eukaryota;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRIO_CALJA
                                                                                                                                                                                                                                                        rion; Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     213 CITQYERESQAYY 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          1 CXTQYXXESXAXY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SIMILARITY: BELONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                   PS00291; PRION_1; 1. PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Metazoa;
                                                                                                                                           230
229
178
180
196
51
                                                                       51
59
67
75
                                                                                                                                                                                                                                                        Glycoprotein;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               STANDARD;
                                          90 5
27639 MW;
                                                                       58
74
82
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chordata;
 80.4%;
61.5%;
                                                                                                                                                                                                                                           GPI-anchor; Repeat; Signal BY SIMILARITY.
Score 41; DB 1; Pred. No. 0.012;
                                                                                                                                      N-LINKED (GLCNAC. . .) (POTENTIAL).
N-LINKED (GLCNAC. . .) (POTENTIAL).
5 X 8 AA TANDEM REPEATS OF P-H-G-G-
                                                                                                                                                                                  REMOVED IN MATURE FORM (BY SIMILARITY).
GPI-ANCHOR (BY SIMILARITY).
BY SIMILARITY.
                                                                                                                                                                                                                            MAJOR PRION PROTEIN.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Craniata; Vertebrata; Euteleostomi;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                PRT;
                                          B2800B60FD5CE664 CRC64;
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             Length 252;
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Gaps

0

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RESULT
PRIO_CE
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              Query Match
 Best Local Similarity
                                                                                                                                                      CARBOHYD
DOMAIN
                                                                                                                                                                                  LIPID
DISULFID
CARBOHYD
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE FROM N.A.

MEDLINE-95139066; PubMed-7837269;

Schatzl H.M., Dacosta M., Taylor L., Cohen F.E., Prusiner S.B.;

"Prion protein gene variation among primates.";

J. Mol. Biol. 245:362-374(1995).

-I- FUNCTION: THE FUNCTION OF PRP IS NOT KNOWN. PRP IS ENCODED IN THE HOST GENOME AND IS EXPRESSED BOTH IN NORMAL AND INFECTED CELLS.

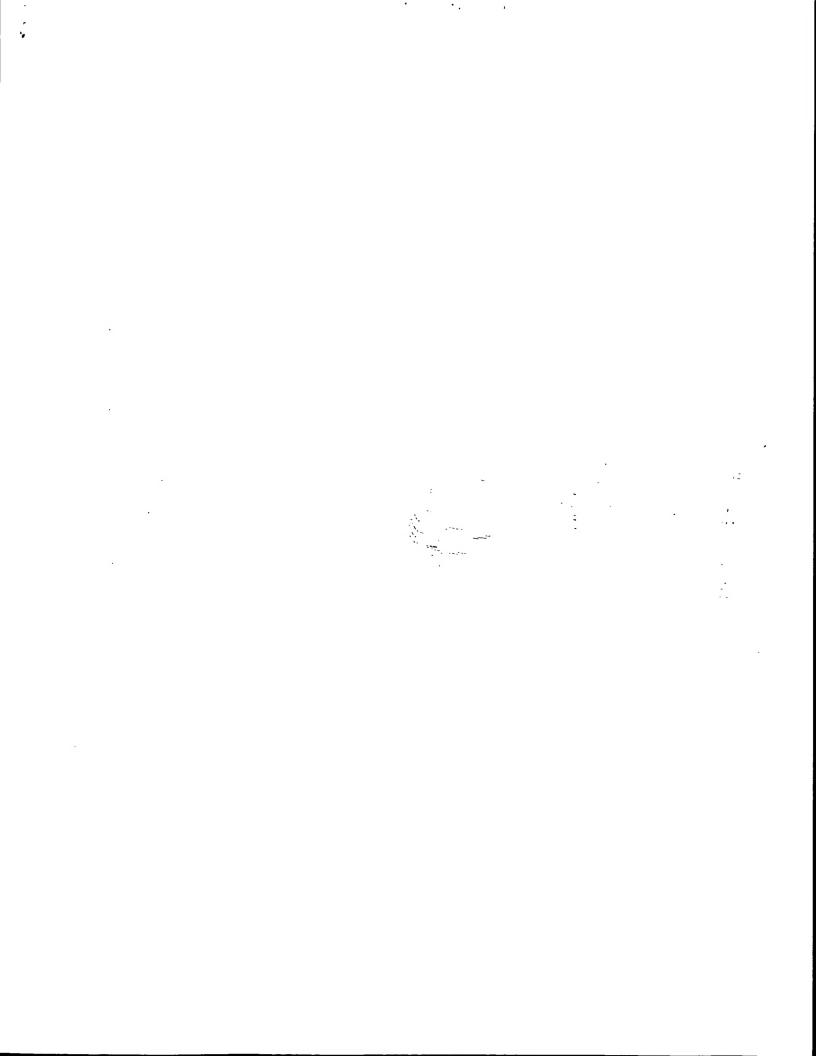
-I- SUBJUNIT: PRP HAS A TENDENCY TO AGGREGATE YIELDING POLYMERS CALLED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cebus apella (Brown-capped capuchin).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Primates; Platyrrhini; Cebidae; Cebinae; Cebus.
NCBI_TaxID=9515;
                                                                                           REPEAT
REPEAT
REPEAT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        This SWISS-PROT entry is copyright. It is produced through a collaboration between the Swiss Institute of Bioinformatics and the EMBL outstation the European Bioinformatics Institute. There are no restrictions on its use by non-profit institutions as long as its content is in no way
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-FEB-1995 (Rel. 31, Created)
01-FEB-1995 (Rel. 31, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
Major prion protein precursor (PrP) (PrP27-30) (PrP33-35C).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CEBAP
                                                                                                                                                                                                                                                                                                                         PRINTS; PR00341; PRION. SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                Pfam; PF00377; prion;
                                                                                                                                                                                                                                                                                                                                                                                                                                               modified and this statement is not removed. entities requires a license agreement (See
                                             SEQUENCE
                                                                                                                                                                                                                                                            SIGNAL
                                                                                                                                                                                                                                                                                          PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                EMBL; U08295; AAC50084.1; -. HSSP; P04156; 1E1G.
                                                                                                                                                                                                                                                                                                                                                                                                                                 or send an email to license@isb-sib.ch).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        P40249;
                                                            REPEAT
                                                                             REPEAT
                                                                                                                                                                                                                                PROPEP
                                                                                                                                                                                                                                                                              Prion;
                                                                                                                                                                                                                                                                                                                                                                     InterPro; IPR000817; Prion.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  213 CITQYEKESQAYY 225
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             1 CXTQYXXESXAXY 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SUBCELLULAR LOCATION: Attached to the membrane by a GPI-anchor. DISEASE: PRP IS FOUND IN HIGH QUANTITY IN THE BRAIN OF HUMANS AND ANIMALS INFECTED WITH THE DEGENERATIVE NEUROLOGICAL DISEASES KURU, CREUTZFELDT-JAKOB DISEASE (CJD), GERSTMANN-STRAUSSLER SYNDROME (GSS), SCRAPIE, BOVINE SPONGIFORM ENCEPHALOPATHY (BSE), TRANSMISSIBLE MINK ENCEPHALOPATHY (TME), ETC.

SIMILARITY: BELLONGS TO THE PRION FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CEBAP
                                                                                                                                                                                                                                                                           Brain;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           .9
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                                          58 1
66 2
74 3
82 4
90 5
27579 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                              license agreement
 80.4%;
                                                                                                                                                                                                                                                              GPI-anchor; Repeat; Signal.
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0;
Score 41; DB 1;
Pred. No. 0.012;
                                                                                                                                                    N-LINKED (GLCNAC. ..) (POTENTIAL).
N-LINKED (GLCNAC. ..) (POTENTIAL).
5 X 8 AA TANDEM REPEATS OF P-H-G-G-W-G-
                                                                                                                                                                                                MAJOR PRION PROTEIN.
REMOVED IN MATURE FORM (BY SIMILARITY).
GPI-ANCHOR (BY SIMILARITY).
BY SIMILARITY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRT;
                                               A2DFCA0AD26D7821 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       252 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                               (See http://www.isb-sib.ch/announce/
                                                                                                                                                                                                                                                                                                                                                                                                                                                               Usage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           <u>ن</u>
              Length 252;
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                                 Matches
213 CITQYERESQAYY 225
                1 CXTQYXXESXAXY 13
                                 8;
                                 Conservative
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                                 Mismatches
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                                 Indels
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Search completed: March 24, 2003, 17:20:17 Job time : 7.85417 secs



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Title:
Perfect score:
Sequence:
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Listing first 45 summaries
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                                                                                                                                                                                                                                       SPTREMBL_21:*
1: sp_archea:*
2: sp_bacteria:
3: sp_fungi:*
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Gapop 10.0 , Gapext 0.5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                       671580 seqs, 206047115 residues
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1 CXTQYXXESXAXY 13
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51
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sp_vertebrate:*
sp_unclassified:*
sp_rvirus:*
sp_bacteriap:*
sp_archeap:*
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sp_bacteria:*
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sp_rodent:*
                                                                                   sp_virus:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

16	15	14	13	12	11	10	9	8	7	6	S	4	w	2		No.	Result	
41	41	41	41	41	41	41	41	41	41	41	41	41	41	42	42	Score		
80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	82.4	82.4	Match	Query	æ
233	227	226	223	220	216	215	214	213	212	202	200	195	195	254	243	Length		
4	6	0	6	6	6	6	σ	6	6	6	6	6	6	σ	11	DB		
P78446	097909	097907	097910	002825	Q9TV00	097904	Q9TV03	Q9TV04	097698	097908	097912	097693	097903	Q9TSF8	P97895	Ħ		
P78446 homo sapien	097909 tragelaphus	097907 gazella sub	097910 hippotragus	002825 odocoileus	Q9tv00 bos taurus		Q9tv03 canis famil	Q9tv04 canis famil	097698 cervus elap	097908 capra nubia	097912 bison bonas	O97693 canis lupus	O97903 addax nasom	Q9tsf8 oryctolagus	P97895 mesocricetu	Description		

44	43	42	41	40	39	38	37	36	35	34	33	32	31	30	29	28	27	26	25	24	23	22	21		19	18	17
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80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4
285	257	256	256	256	256	256	256	256	256	256	256	256	256	256	254	254	254	253	253	253	253	248	246	245	245	240	235
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Q9mzu6 075942	046593	Q8spv4	Q8spv5	08spv6	Q8spv7	046648	062670	002841	Q95m08	095n12	Q9mzu8	Q9ttu5	Q9tu07	Q9tv01	08vhv6	09qyt	Q9z0t4	Q9z0t5	Q8tbg0	Q96e70	Q9up19	Q8vhv!	060489	Q9mzu7	Q15216	Q8vhv4	097695
antilocapra homo sapien			capra		capra hircu	capra hircu	cervus elap	odocoileus	budorcas ta	ovis aries	odocoileus	ovis aries	ovis aries	a	6 apodemus sy			Ç,			homo sapien	Q8vhv5 clethrionom	homo sapien	odocoileus	⇁	microtu	giraffa cam

ALIGNMENTS

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SO	ΚW	DR	DR	DR	DR	DR	DR	DR	DR	DR	RL	RΤ	RA	RA	RΑ	RX	RΡ	RN	RL	RT	₽₽	RX	RΡ	RN	×	8	გ	8	S	DE	ΡŢ	ဌ	Ŋ	AC	ID	P97895	RESULT
NON_TER 1 1 SEQUENCE 243 AA; 26643 MW; 4F53612BBFF240F9 CRC64;			E; PS00291		; PRION	Pfam; PF00377; prion; 1.	InterPro; IPR000817; Prion.	HSSP; P04273; 1B10.	K02234; AAA37093.1;	EMBL; M37381; AAA37090.1;	1 40:735-746(1985).	"A cellular gene encodes scrapie PrP 27-30 protein.";	siner S.B., Weissmann C.;	P., Teplów D.B., Hood	Oesch B., Westaway D., Waelchli M., McKinley M.P., Kent S.B.,	MEDLINE=85176927; PubMed=2859120;	SEQUENCE OF 79-223 FROM N.A.	[2]	Int. Rev. Neurobiol. 28:1-57(1986).	"Biology and structure of scrapie prions.";	McKinley M.P., Prusiner S.B.;	MEDLINE-87108309; PubMed-3100471;	SEQUENCE FROM N.A.	[1]	NCBI_TaxID=10036;		Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Cricetinae;	Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;	uratus (Gold	Scrapie prion (PRP 27-30) (PRP 27-30 protein) (Fragment).	rEMBLrel. 17	(TrEMBLrel.	(TrEMBLrel.	P97895;	P97895 PRELIMINARY; PRT; 243 AA.	395	JLT 1

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Best Local
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Pfam; PF00377; Prion; 1.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
PROSITE; PS000291; PRION_1; 1.
PROSITE; PS00076; PRION_2; 1.
SEQUENCE 254 AA; 27546 MM;
                                                                                                                                                                                                01-MAY-1999
01-MAY-1999
01-JUN-2002
MEDLINE-99303687; PubMed=10373359; Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schatzl H.M.; Renalysis of 27 meanmalian and 9 avian Prps reveals high conservation of flexible regions of the prion protein."; J. Mol. Biol. 289:1163-1178(1999).
                                                                                                                                                                                                                                     097903;
                                                                                                                                   Addax nasomaculatus.
Eukaryota; Metazoa; Chordata; Craniata; Vertebra;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           immune response to the prion protein immunogen.";
Neurovirol. 5:401-413(1999)
EMBL; AF015603; AAD01554.1;
HSSP; P10279; IDWY
HSSP; P10279; IDWY
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     PRP.
Oryctolagus cuniculus (Rabbit).
Curotolagus cuniculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Eukaryota; Metazoa; Chordata; Craniata; Oryctolagus.
                                                                           TISSUE-PBL;
                                                                                     SEQUENCE FROM N.A.
                                                                                                             Bovidae; Hippotraginae;
NCBI_TaxID=59515;
                                                                                                                                                                                      Prion protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE FROM N.A. MEDLINE-99391519; PubMed-10463862;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAY-2000 (TrEMBLrel.
01-MAY-2000 (TrEMBLrel.
01-JUN-2002 (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Rubenstein R., Kascsak R.J., Papini M., Kascsak R., Carp R.I.,
LaPauci G., Meloen R., Langeveld J.;
"Immune surveillance and antigen conformation determines humoral
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Q9TSF8
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NCBI_TaxID=9986;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Prion protein.
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                                                                                                                                                                                                                                                                                                            CITOYQQESQAAY 227
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CTTQYQKESQAYY 215
                                                                                                                                                                                                                                                                                                                                                             Similarity
8; Conserv
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8; Conser
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                                                                                                                                                                                                                                                                                                                                                                       82.4%;
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13,
21,
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Last sequence update)
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                      Score 42; DB 6;
Pred. No. 0.017;
                                                                                                                                                                                                                                                  PRT;
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                                                                                                                                                                                                                                                  195
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                                                                                                                                  Vertebrata; Euteleostomi;
minantia; Pecora; Bovoidea
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0.017;
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                                                                                                                                    Bovoidea;
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RESULT 4
097693
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Best Local Similarity binches 8; Conservative
                                                                                                                                                                                                                                                                                                                                Matches
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Best Local
097912 PRELIMINARY;
097912;
01-MAY-1999 (TREMBLrel. 10, C
01-MAY-1999 (TREMBLrel. 21, L
01-JUN-2002 (TREMBLrel. 21, L
Prion protein (Fragment).
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PRINTS; PR00334; KININGEN.
PRINTS; PR00334; PRION.
SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
NON_TER 195 195
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PROSITE; PS00706; PRION_2;
NON_TER 1 1
NON_TER 195 195
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Schwarz T.F., Werner T., Schatzl H.M.;
"Analysis of 27 mammalian and 9 avian PrPs
of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAY-1999 (TrEMBLrel. 10, 01-MAY-1999 (TrEMBLrel. 10, 01-DEC-2001 (TrEMBLrel. 19,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             InterPro; IPR000817; Prion. Pfam; PF00377; Prion; 1. PRINTS; PR00341; PRION. SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae;
NCBI_TaxID=9612;
                                                                                                                                                                                                                                                                                                                                                                                                                     SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    EMBL; AF113939; AAD12063.1;
HSSP; P04925; lAG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
MEDLINE-99303687; PubMed-10373359;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Canis lupus (Gray wolf).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Prion protein
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InterPro; IPR000817; Prion.
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                                                                                                                                                                                                                                                                                  1 CXTOYXXESXAXY 13
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8; Conserv
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195 AA;
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195 AA;
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; 21321 MW;
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21097 MW; 9D18E4EB9AA5D031 CRC64;
                                                                                                                                                                                                                                                                                                                                                80.4%;
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                                Created)
Last sequence update)
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                                                                                                                                                                                                                                                                                                                           Score 41; DB
Pred. No. 0.02
0; Mismatches
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0.023;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Euteleostomi;
Canis.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gilch
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Best Local S
Matches 8
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PRINTS; PR00334; KININGGEN.
PRINTS; PR00341; PRION.
SMART; SM00086; PAC; 1.

SMART; SM00157; PRP; 1.

PROSITE; PS00291; PRION_1; 1.

PROSITE; PS00706; PRION_2; 1.
Interpro: IPR00239; Kininogen.
Interpro: IPR002817; Prion.
Pfam; PP00377; prion; 1.
PRINTS; PR00334; KININCEN.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP: 1.
PROSITE; PS00706; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
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Schwarz G.F., Werner T., Schatzl H.M.;
"Analysis of 27 mammalian and 9 avian PrPs reveals high of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalla; Euthorla; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;

Bovidae; Bovinae; Bison.
                                                                                                                                                                                                               Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Schwarz T.F., Werner T., Schatzl H.M.
"Analysis of 27 mammalian and 9 avian PrPs reveals high of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
EMBL; AFIJ7319; AAD19990.1;
EMBL; AFIJ7319; AAD19990.1;
HSSP; P10279; IDWY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoldea Bovidae; Caprinae; Capra.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            01-MAY-1999 (TrEMBLrel. 10, 01-MAY-1999 (TrEMBLrel. 10, 01-JUN-2002 (TrEMBLrel. 21,
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Capra nubiana (Nubian ibex).
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                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE=99303687; PubMed=10373359;
                                                                                                                                                                                                                                                                                                                                                                                                                                 TISSUE-PBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Prion
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InterPro; IPR001610; PAC.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          182
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200 AA;
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Last annotation update)
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Pred. No. 0.024;
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                                                                                                                                                                                                                                                                                                                                                                               Gilch S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Bovoidea;
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Best Local
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O97698;
O1-MAY-1999 (TrEMBLrel. 10, C
O1-MAY-1999 (TrEMBLrel. 10, I
O1-JUN-2002 (TrEMBLrel. 21, I
Prion protein (Fragment).
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InterPro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
PRINTS; PR00334; KININOGEN.
PRINTS; PR00341; PRION.
PRINTS; SM00157; PRP; 1.
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NON_TER
SEQUENCE
        Q9TV04 PRELIMINARY;
Q9TV04;
01-MAY-2000 (TIEMBLrel. 13,
01-MAY-2000 (TIEMBLrel. 13,
01-DEC-2001 (TIEMBLrel. 19,
                                                                                                                                                                                                                    PROSITE; PS00291; PRION_1;
PROSITE; PS00706; PRION_2;
NON_TER 1 1
NON_TER 212 212
SEQUENCE 212 AA; 23032 M
                                                                                                                                                                                                                                                                                                                                                                  MEDIINE=99303687; pubMed=10373359; Wopfner F., Weldenhofer G., Schmelder R., von Brunn A., Gilch S., Schwarz T., Weldenhofer G., Schmelder R., von Brunn A., Gilch S., Schwarz T., F., Werner T., Schatzl H.M.; "Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein."; of flexible regions of the prion protein."; J. Mol. Biol. 289:1163-1178(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SEQUENCE FROM N.A.
MEDLINE-97317556; PubMed-9174569;
Schatzl H.M., Wopfner F., Gilch S.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Eukaryota; Metazoa;
Mammalia; Eutheria;
Cervidae; Cervinae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cervus elaphus canadensis (wapiti).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Cervoide
Prion
                                                                                                                                                                                                                                                                                                                                                         HSSP; P10279; 1DWY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          animals?"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   SCHRIZI H.M., WOPINET F., Gilch S., von Brunn
"Is codon 129 of prion protein polymorphic in
animalia".
                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NCBI_TaxID=9861;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   189 CITOYORESOAYY 201
                                                                                                                       181 CITQYQRESEAYY 193
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ancet
                                                                                                                                                                     Local Similarity
nes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           1 CXTQYXXESXAXY
                                                                                                                                             1 CXTQYXXESXAXY 13
protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                349:1603-1604(1997).
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8; Conserv
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202
202
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202
21949 MW;
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23032 MW; 57E58ABDF5E2A1B5 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Cervus.
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                                                                                                                                                                                                                                                                                                                                          Kininogen.
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Last annotation updat
          Last sequence update)
Last annotation update)
                                   Created)
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                                                                                                                                                                     Score 41; DB 6;
Pred. No. 0.025;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 41; DB 6; Pred. No. 0.024;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    DB0634A43B4DB77F CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Ruminantia; Pecora; Cervoidea;
                                                           213
                                                                                                                                                                                            DB 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         A
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                                                                                                                                                                                            Length 212;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Jager G.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        beings but not in
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RESULT 9
Q9TV03
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         Matches
                                 Query Match
Best Local
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Best Local Similarity
Matches 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              InterPro: IPRO00817; Prion.
Pfam; PR00377; Prion; 1.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                   Wopfner F., Weldenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schwatzl H.M., Schwarz G. T. Werner T., Schwatzl H.M., "Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein.";
J. Mol. Biol. 289:1103-1178(1999).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           01-MAY-2000
01-MAY-2000
01-DEC-2001
                                                                                                                                                                                             PROSITE;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CANIS familiaris (Dog).
Canis familiaris (Dog).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata;
Eukaryota; Metazoa; Chordata; Cranidae; Canidae;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Q9TV03;
                                                                                                                                                                                                                                      InterPro; IPRO00817; Prion.
Pfam; PF00377; Prion; 1.
PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Eukaryota; Metazoa; Chordata; (Mammalia; Eutheria; Carnivora; NCBI_TaxID=9615;
                                                                                                       SEQUENCE
                                                                                                                                                                                                                                                                                                                                  EMBL; AF113938; AAD12062.1; HSSP; P04925; 1AG2.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       MEDLINE-99303687; PubMed-10373359;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schatzl H.M.;
"Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein.";
J. Mol. 18101 289:1163-1178 (1999)
                                                                                                                          NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Prion protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            EMBL; AF113937; AAI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SEQUENCE FROM N.A. MEDLINE-99303687; PubMed-10373359;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Carnivora; Fissipedia; Canidae; Canis.
                                                                                                                                                  /ARIANT
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        177
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         Local Similarity
nes 8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 1 CXTQYXXESXAXY 13
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   CVTQYQKESEAYY 189
                                                                                                                                                                                         PS00291; PRION_1; 1. PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     213
213 AA;
                                                                                                    214 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    (TrEMBLrel. 13, Created)
(TrEMBLrel. 13, Last sequence update)
(TrEMBLrel. 19, Last annotation updat
         Conservative
                                                                                                                        214
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRELIMINARY;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    AAD12061.1;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     213
22997 MW;
                                                                                                23167 MW;
                            80.4%;
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       0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
    Score 41; DB 6; Pred. No. 0.026; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Score 41; DB 6;
Pred. No. 0.026;
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v
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                                                                                                551B7669ABD4C6DF CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    214 AA
                                                  6.
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                                                  Length 214;
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    Indels
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    0;
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Gaps
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Q9TV00
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InterPro; IPR001610; PAC.
InterPro; IPR000817; Prion.
InterPro; IPR000817; Prion.
Pfam; PF00377; prion; 1.
PRINTS; PR00374; KININOGEN.
PRINTS; PR00341; PRION.
SMART; SM00086; PAC; 1.
SMARF; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-MAY-1999
01-MAY-1999
01-JUN-2002
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Q9TV00
Q9TV00;
Q1TV00;
Q1-MAY-2000 (TrEMBLrel. 13,
Q1-MAY-2000 (TrEMBLrel. 13,
Q1-JUN-2002 (TrEMBLrel. 21,
MEDLINE-99303687; PubMed-10373359; Wopfner F., Weidenhofer G., Schneider R., Schwarz T.F., Werner T., Schatzl H.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       NON_TER
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Wopfner F., Weidenbofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schatzl H.M.;

"Analysis of 27 mammalian and 9 avian PrPs reveals high conservation of flexible regions of the prion protein.";

J. Mol. Biol. 289:1163-1178(1999).

EMBL; AF117310; AAD19981.1;

TELEPTIC TO BENEFIT OF THE PROPERTY OF THE PROPER
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                                                                                                                                                                                                                                                Bos taurus (Bovine).
Eukaryota; Metazoa; Chordata;
Mammalia; Eutheria; Cetartioda
Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                                                                                                                                                                                                   PRP.
                                                                                                               SEQUENCE FROM N.A.
TISSUE-PBL;
                                                                                                                                                                                                                   NCBI_TaxID=9913;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Bos javanicus (Wild banteng).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
                                                                                                                                                                                                                                                                                                                                                                                                                            Prion protein (Fragment).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-99303687; PubMed-10373359;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Prion protein (Fragment)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID-9906;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      190 CITOYORESOAYY 202
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         177
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Local
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 CXTQYXXESXAXY 13
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Similarity
8; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   215
215 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (TrEMBLrel. (TrEMBLrel.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (TrEMBLrel.
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                                                                                                                                                                                                                                                                            Chordata; Craniata; Vertebrata; Euteleostomi; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   23182 MW;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              80.4%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     10,
10,
21,
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Created)
Last sequence update)
Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0;
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Last sequence up
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Score 41; DB 6;
Pred. No. 0.026;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    PRT;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        O'Rourke K.I., Spraker T.R., Miller M.W., Williams E.S.; "Three alleles of the prion protein gene in mule deer (Odocolleus hemionus) with chronic wasting disease."; Submitted (APR-1997) to the EMBL/GenBank/DDBJ databases. EMBL; U97331; AAB61573.1; -.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalla; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Cervoide
Cervidae; Odocoileinae; Odocoileus.
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SMART; SM00157; PRF; 1.
PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
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SEQUENCE
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PRINTS; PR00341; PRION.
SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION. 1;
PROSITE; PS00706; PRION_2;
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J. Mol. Biol. 289:1163-1178(1999).
EMBL; AFI17327; AAD19998.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           TISSUE-BRAIN;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NCBI_TaxID=9877;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Prion protein (Fragment).
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InterPro; IPR000817; Prion.
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Pred. No. 0.026;
                                                                                                                                                                                                                                                                                                                                                                           Score 41; DB 6;
Pred. No. 0.027;
       PRT;
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Best Local Similarity
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                               of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
EMBL; AF117313; AAD19984.1; -.
HSSP; P10279; 1DWY.
Interpro; IPR002395; Kininogen.
Interpro; IPR000817; Prion.
Pfam; PF00377; Prion; 1.
PRINTS; PR00334; KININOGEN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hippotragus niger (Sable antelope).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea
Bovidae; Hippotraginae; Hippotragus.
NCBI_TaxID-37189;
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01-MAY-1999 (TrEMBLrel. 10,
01-MAY-1999 (TrEMBLrel. 10,
01-JUN-2002 (TrEMBLrel. 21,
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SMART; SM00157; PRP; 1.
PROSITE; PS00291; PRION_1;
PROSITE; PS00706; PRION_2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gazella subgutturosa (Goitred gazelle).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
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                                                                                                                                                                                                                                                                                                                                        MEDLINE-99303687; PubMed-10373359; Wopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S., Schwarz T.F., Werner T., Schwazl H.M.; Schwarz T.F., Warner T., Schwazl H.M.; The schwarz T.F. werner T., Schwazl H.M.; The schwarz T.F. werner T., Schwarz Schwarz T. werner T., Werner T., Schwarz T. werner 
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PRINTS; PR00334; KININOGEN.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           NCBI_TaxID=59529;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Prion protein (Fragment).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  190 CITQYQRESQAYY 202
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223 AA;
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24172 MW;
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Last sequence update)
Last annotation update)
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Pred. No. 0.027;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  226 AA
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RESULT 15
097909
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Search completed: March 24, 2003, 17:22:11 Job time: 28.4375 secs
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                                                                                                                                                                                                                                           Query Match 80.4%; Score 41; DB 6; Length 227; Best Local Similarity 61.5%; Pred. No. 0.027; Matches 8; Conservative 0; Mismatches 5; Indels
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Best Local Similarity 61.5%;
Matches 8; Conservative (
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PROSITE; PS00291; PRION_1; 1.
PROSITE; PS00706; PRION_2; 1.
NON_TER 1 209 209
VARIANT 209 206
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SEQUENCE
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MEDLINE-9303687; PubMed-10373359;
Mopfner F., Weidenhofer G., Schneider R., von Brunn A., Gilch S.,
Schwarz T.F., Werner T., Schatzl H.M.;
Schwarz T.F., Werner T., Schatzl H.M.;
Analysis of 27 mammalian and 9 avian Prps reveals high conservation of flexible regions of the prion protein.";
J. Mol. Biol. 289:1163-1178(1999).
EMBL; API17321; AnaD1992.1; -.
EMBL; API17321; AnaD1992.1; -.
HSSP; P10279; IDWY.
InterPro; IPR002395; Kininogen.
InterPro; IPR003395; Kininogen.
Pfam; PF00377; prion; 1.
PRINTS; PR00334; KININOGEN.
PRINTS; PR00334; KININOGEN.
SMART. SMONIS7; DED. 1
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O97909; O1-MAY-1999 (TrEMBLrel. 10, Created)
O1-MAY-1999 (TrEMBLrel. 10, Last sequence update)
O1-JUN-2002 (TrEMBLrel. 21, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Tragelaphus angasii (Nyala).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Tragelaphus.
NCBI_TaxID=66437;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE FROM N.A.
TISSUE-PBL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   PROSITE; PS00291; PRION_1; 1. PROSITE; PS00706; PRION_2; 1.
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                                                                                                                         201 CITOYRRESOAYY 213
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  SMART; SM00157; PRP; 1.
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226 226 226
226 AA; 24384 MW; D845E27B219ABD2F CRC64;
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24651 MW; 99FD0BAF0B6A0077 CRC64;
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Pred. No. 0.027;
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Minimum DB
Maximum DB
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Perfect score:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Database
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  Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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seq length: 2000000000
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1: /cgn2_6/ptodata/2/
2: /cgn2_6/ptodata/2/
3: /cgn2_6/ptodata/2/
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51
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 Maximum Match 100%
Listing first 45 summaries
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/cgn2_6/ptodata/2/1aa/5B_COMB.pep:*
/cgn2_6/ptodata/2/1aa/6A_COMB.pep:*
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/cgn2_6/ptodata/2/1aa/backfiles1.pep:*
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US-09-076-721-48
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US-08-244-701B-35
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US-08-244-701B-37
US-09-076-721-35
US-09-076-721-35
US-09-076-721-62
US-09-076-721-62
US-09-353-348-7
US-08-542-188-2
US-08-509-261R-2
US-08-660-626-8
US-08-692-892-2
US-08-162R-22
US-09-031-168R-22
US-09-031-168R-22
US-09-031-168R-22
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US-09-128-450-26
US-09-823-494-26
US-08-244-7018-48
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Sequence 26, Appl
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	US-08-242-188-1 US-08-509-261A-1 US-08-660-626-7
1, 21, 12, 13, 14, 14, 14, 14, 14, 14, 14, 14, 14, 14	Sequence 1, Appl Sequence 1, Appl Sequence 7, Appl

ALIGNMENTS

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; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-556-823-2
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Best Local Similarity
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                                                                                                                                                                              TELEFAX: (415) 854-0875
INFORMATION FOR SEQ ID NO: 2:
SEQUENCE CHARACTERISTICS:
LENGTH: 142 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                  COMPUTER READABLE FORM:

MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: ASCIII
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/556,823
FILING DATE:
CLASSIFICATION: 530
ATTORNEY/AGENT INFORMATION:
NAME: Valeta Gregg
REGISTRATION NUMBER: 07532/00:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (415) 322-5070
TELEPHONE: (415) 322-5070
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TITLE OF INVENTION: FOIR TITLE OF INVENTION: NUMBER OF SEQUENCES: 10 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  APPLICANT:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ADDRESSEE: Fish & Richardson
STREET: 2200 Sand Hill Road, Suite 100
CITY: Menlo Park
STATE: California
COUNTRY: USA
                                   1 CXTQYXXESXAXY 13
CTTQYQKESQAYY 137
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Fred E. Cohen
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                                                                       Conservative
                                                                                       82.4%;
61.5%;
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Pred. No. 0.014;
                                                                       Mismatches
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RESULT 4
US-08-244-701B-48
; Sequence 48, Application US/08244701B
Parent No. 5773572
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; TYPE: PRT
; ORGANISM: Hamster sp.
US-09-823-494-26
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US-09-128-450-26
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; TYPE: PRT
; ORGANISM: Hamster sp.
US-09-128-450-26
                                                                                                                                                                                                                                                                                                                                                             PRIOR FILING DATE: 1998-08-03
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PATENTIN VET. 2.0
SEQ ID NO 26
LENGTH: 254
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Patent No.
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APPLICANT: Chesebro, Bruce W
APPLICANT: Chashry, Byron W
APPLICANT: Chabry, Joelle
APPLICANT: Chabry, Joelle
APPLICANT: Priols, Susette
TITLE OF INVENTION: Inhibitors of Formation of Protease Resistant Prion
TITLE OF INVENTION: Protein
FILE REFERENCE: 50121
                                                                                                                                                                                                                                                Matches
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CURRENT FILING DATE: 1998-08-03
NUMBER OF SEQ ID NOS: 29
SOFTWARE: PATENTIN Ver. 2.0
SEQ ID NO 26
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                GENERAL INFORMATION:
APPLICANT: Fishle
APPLICANT: Robson
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APPLICANT: Caughey, Byron W
APPLICANT: Caughey, Byron W
APPLICANT: Chabry, Joselle
APPLICANT: P101a, Susette
TITLE OF INVENTION: Inhibitors of Formation of Protease Resistant Prion
FILE REFERENCE: 50121
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                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/823,494
CURRENT FILING DATE: 2001-03-30
PRIOR APPLICATION NUMBER: 09/128,450
   APPLICANT:
                                                                                                                                                                          214 CTTQYQKESQAYY 226
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Local Similarity 61.5%;
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Local Similarity 61.5%;
es 8; Conservative
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635561
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Fishleigh, Robert V. Robson, Barry Mee, Roger P.
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Pred. No. 0.
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Pred. No. 0
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0.027;
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US-09-076-721-48
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICATION NUMBER: US/08/244
FILING DATE: 02-JUN-1994
CLASSIFICATION: 436
ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 8080
TELECOMMUNICATION INFORMATION:
                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                              APPLICANT:
APPLICANT:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                             SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA:
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COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0.
CURRENT APPLICATION DATA:
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                                                                                                                                                                                                                                                                                 NUMBER OF SEQUENCES:
                                                                                                                                                                                                                                                                                                 TITLE OF INVENTION:
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NUMBER OF SEQUENCES:
                             FILING DATE:
CLASSIFICATION:
                                                                                        SOFTWARE:
                                                                                                                                                                                                     STATE: New York
                                                             APPLICATION NUMBER:
                                                                                                                                                                                                                                   STREET:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Local Similarity
les 8; Conserv
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                                                                                                                                                                                      COUNTRY:
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o. 6379905
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1155 Avenue of the Americas
                                                                                                                                                                                                                                                                                                          Robson, Barry
Mee, Roger P.
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(212) 869-8864/9741
                                                                                                                                                                                                                                                                                                                                           Fishleigh, Robert V.
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                                                             US/09/076,721
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US 08/244,701

ATTORNEY/AGENT INFORMATION:

Fanucci, Allan A.

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Query Match
Best Local Similarity
Thehes 8; Conserve
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; TYPE: PRT
; ORGANISM: Ovis aries
US-09-353-348-3
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                                                                                                                                                                                                            Sequence 33, Application US/08244701B Patent No. 5773572
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FILE REFERENCE: OROUTKE
CURRENT APPLICATION NUMBER: US/09/353,348
CURRENT FILING DATE: 1999-07-15
NUMBER OF SEQ ID NOS: 11
                                                                                                                                                                                                GENERAL INFORMATION:
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TELECOMMUNICATION INFORMATION:
TELEPHONE: (212) 790-9090
                                                                                                   APPLICANT: Mee, Roger P.
TITLE OF INVENTION: Eragments of Prion Proteins
NUMBER OF SEQUENCES: 67
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: peptide
                                                                                                                                                                                APPLICANT:
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COMPUTER READABLE FORM:
           COUNTRY: U.
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les 8; Conserve
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                                                          New York
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VENTION: Monoclonal Antibodies and Antibody Cocktail

VENTION: Detection of Prion Protein as an Indication

VENTION: Transmissible Spongiform Encephalopathies
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                              U.S.A.
                                                                                                                                                                Fishleigh, Robert V. Robson, Barry
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                                                                                        Pennie & Edmonds
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Pred. No. 0.
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TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                GENERAL INFORMATION:
                                                                 MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
NAME: Fanucci, Allan A.
REGISTRATION NUMBER: 30,256
                                                                                                                                         COMPUTER READABLE FORM: MEDIUM TYPE: Floppy
                                                                                                                                                                                                                                                                                                                                        APPLICANT: Fishleigh, RO APPLICANT: Robson, Barry APPLICANT: Mee, Roger P.
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CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                      NUMBER OF SEQUENCES:
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ATTORNEY/AGENT INFORMATION:
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OTHER INFORMATION:
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TYPE: amino acid
STRANDEDNESS: single
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               APPLICATION NUMBER: FILING DATE: 02-JUN CLASSIFICATION: 436
                                                                                                                                                                                                                CITY: New York
STATE: New Yor
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mes 8; Conserv
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                                                                                                                                                                                ZIP: 10036
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Y: U.S.A.
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                                                                                                                                                                                                                                                                                                        Fragments of Prion Proteins 67
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 80.4%;
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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Pred. No. 0.0033;
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US-08-244-701B-37
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                                  INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Fishleign, north applicant: Robson, Barry APPLICANT: Mee, Roger P. TITLE OF INVENTION: Fragment Frag
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                                                                                                                                                                      ATTORNEY/AGENT INFORMATION:
NAME: FANUCCI, Allan A.
REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 801
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                             SOFTMARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/244,701B
FILING DATE: 02-UUN-1994
CLASSIFICATION: 436
SEQUENCE CHARACTERISTICS:
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REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 800
TELECOMMUNICATION INFORMATION:
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                                                                                                  TELEFHONE: (212) 790-9090
TELEFAX: (212) 869-8864/9741
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OPERATING SYSTEM: PC-DOS/MS-DOS
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LOCATION:
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TYPE: amino acid
STRANDEDNESS: single
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                                                                66141 PENNIE
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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Best Local Similarity
"hthes 8; Conserve
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                                                                                  TELEFAX: (212) 869-886
TELEX: 66141 PENNIE
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
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APPLICANT:
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TITLE OF INVENTION: Fragments of Prion Proteins
NUMBER OF SEQUENCES:
CORRESPONDENCE DADRESS:
MOLECULE TYPE:
FEATURE:
                                                                                                                                                                     REFERENCE/DOCKET NUMBER: 80 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                              CLASSIFICATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
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CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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                              TOPOLOGY:
                                          STRANDEDNESS:
                                                                                                                                                                                                    NAME: Fanucci, Allan A
REGISTRATION NUMBER: 3
                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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OTHER INFORMATION:
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LOCATION:
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                                                                       22 amino acids
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                                                                                                                                        (212) 869-8864/9741
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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MEDIOM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
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REFERENCE/DOCKET NUMBER: 80
TELECOMMUNICATION INFORMATION:
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PRIOR APPLICATION DATA:
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                                                                                                                                LENGTH: 22 amino acids
TYPE: amino acid
STRANDEDNESS: single
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1155 Avenue of the Americas
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                                                                                                                                                                                                                                    (212) 869-8864/9741
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                                                                                                                  SS: single
linear
                                                                                                                                                                                                                                                   (212) 790-9090
                                                                                                   peptide
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
/label= X
/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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Pred. No. 0.0033;
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GENERAL INFORMATION:
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                                                                                      FEATURE:
                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
LENGTH: 22 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SOFTWARE: PatentIn Release #1.0, CURRENT APPLICATION DATA:
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MEDIUM TYPE: Floppy disk
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LOCATION:
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                         OTHER INFORMATION: OTHER INFORMATION:
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STATE: New Yor
                                                                                                                 OTHER INFORMATION: OTHER INFORMATION:
                                                                                                                                               NAME/KEY:
                                                                                                                                                                                                           TOPOLOGY:
                                                                                                                                                                                                                                                                                                                              TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                            NAME: Fanucci, Allan A. REGISTRATION NUMBER: 30, REFERENCE/DOCKET NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE:
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            INFORMATION:
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/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
/label= Y
/note= "Y may be absent or present independently
of X and denotes one or more amino acid(s)"
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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                                                                                                                                                                                                                                                                                                                                     SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
TYPE: amino acid
STRANDEDNESS: single
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MEDIUM TYPE: Floppy
FEATURE
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CORRESPONDENCE ADDRESS:
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    LOCATION: 9
OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Fanucc1, Allan A.
REGISTRATION NUMBER: 30,256
REFERENCE/DOCKET NUMBER: 80
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
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(212) 869-8864/9741
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                                                                               /label= x
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/note= "X may be absent or present independently
of Y and denotes one or more amino acid(s)"
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/note= "X = either Ile, Thr or Val"
        /label= X
/note= "X = Arg or Lys"
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Best Local Similarity
Matches 11; Conserv
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APPLICANT: Fishle
APPLICANT: Robson
APPLICANT: Mee, Ro
PRIOR APPLICATION DATA:
                                         CURRENT APPLICATION DATA:
APPLICATION NUMBER: US,
                                                                                                                         COMPUTER READABLE FORM:
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NUMBER OF SEQUENCES:
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LOCATION:
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LOCATION:
                                                                                COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                            COUNTRY: U.S.A. ZIP: 10036
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                              FILING DATE
                                                                                                            MEDIUM TYPE: Floppy disk
                                                                                                                                                                       STATE:
                                                                                                                                                                                   CITY: New York
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                CLASSIFICATION:
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1155 Avenue of the Americas
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                                                                  PatentIn Release #1.0, Version #1.25
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/note= "X = Asp or Gln"
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/note= "X =
                                          US/09/076,721
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Pred. No. 0.0036;
"Hammatches 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       either Ala, Thr, Met or Val"
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APPLICATION NUMBER:

US 08/244,701

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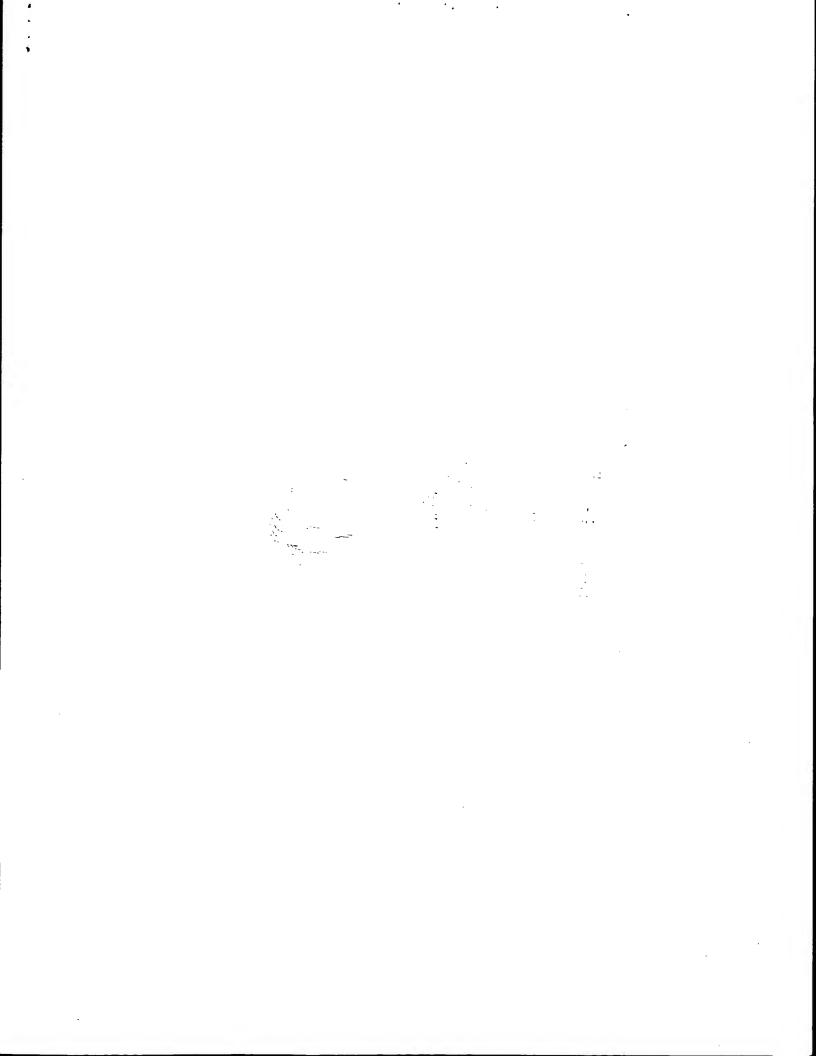
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INFORMATION FOR SEQ ID NO: 62:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS:
LENGTH: 24 amino acids
TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           REFERENCE/DOCKET NUMBER: 8080-007
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        MOLECULE TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
                                                                                                                                                                                                                                                                               LOCATION: 20
OTHER INFORMATION: /label= X
OTHER INFORMATION: /note= "X =
                                                                                                                                                                                                                                                                                                                                                                            LOCATION: 19
OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: Modified-site LOCATION: 17
OTHER INFORMATION: /labe OTHER INFORMATION: /not
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OTHER INFORMATION:
OTHER INFORMATION:
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OTHER INFORMATION: /note- "X -
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OTHER INFORMATION: /note= "X may be absent or present independently
OTHER INFORMATION: of Y and denotes one or more amino acid(s)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NAME: Fanucci, Allan A. REGISTRATION NUMBER: 30,256
                                                                                             NAME/KEY: modified-site LOCATION: 23
OTHER INFORMATION: /labe OTHER INFORMATION: /note
                                                                                                                                                                                        NAME/KEY: modified-site LOCATION: 22 OTHER INFORMATION: /labe OTHER INFORMATION: /note
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OTHER INFORMATION: /note= "X =
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TELEX: 66141 PENNIE
                                                                                                                                                                                                                                                                                                                                                                                                                       NAME/KEY: Modified-site 
LOCATION: 19
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/note- "x -
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/note- "X =
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/note- "X -
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/note= "X = Gly or Arg"
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/note- "X -
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/note- "X -
/label- Y note- "Y may be absent or present independently
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                                                                                                 either Ala, Thr, Met or Val"
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; OTHER INFORMATION: US-09-076-721-62
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US-09-353-348-7
Search completed: March 24, 2003, 17:23:05
Job time : 12.4583 secs
                                                                                                                                                                                              ; ORGANISM: Ovis aries
US-09-353-348-7
                                                                                                                                                                                                                           CURRENT FILING DATE: 1999-07-15
NUMBER OF SEQ ID NOS: 11
SOFTWARE: PatentIn Ver. 2.1
SEQ ID NO 7
LENGTH: 26
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 7, Application US/09353348 Patent No. 6261790 GENERAL INFORMATION:
                                                                                                                         Query Match 80.4%;
Best Local Similarity 61.5%;
Matches 8; Conservative
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                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION:
TITLE OF INVENTION:
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                             CURRENT APPLICATION NUMBER: US/09/353,348
                                                                                                                                                                                                                                                                                                                                                FILE REFERENCE:
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nes 11; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 CXTQYXXESXAXY 13
                                                             5 CITQYQRESQAYY 17
                                                                                            1 CXTQYXXESXAXY 13
                                                                                                                                                                                                                                                                                                                                                       O'Rourke, Katherine I.

VENTION: Monoclonal Antibodies and Antibody Cocktail
VENTION: Detection of Prion Protein as an Indication
VENTION: Transmissible Spongiform Encephalopathies
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
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                                                                                                                           Score 41; DB 4; LC Pred. No. 0.0039; ...
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Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 4;
0.0036;
                                                                                                                                                            DB 4; Length 26;
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Title:
Perfect score:
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Maximum Match 100%
Listing first 45 summaries
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Minimum DB seq length: 0
Maximum DB seq length: 2000000000
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                                                                                                                                                                                                                                                                                                                                                                                                  Database :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Searched:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Scoring table:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       US-09-508-828B-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       March 24, 2003, 17:23:11; Search time 11.6458 Seconds (without alignments) 59.679 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                221153 seqs, 53462247 residues
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        GenCore version 5.1.4_p5_4578 Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                   /cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
/cgn2_6/ptodata/1/pubpaa/PCT_NEW_PUB.pep:*
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed. and is derived by analysis of the total score distribution.

SUMMARIES

19	18	17	16	15	14	13	12	11	10	9	8	7	6	5	4	u	2	1	Result
41	41	41	41	41	41	41	41	41	41	41	41	41	41	41	41	41	41	42	Score
80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	80.4	82.4	Query Match Length
256	256	256	256	256	255	254	254	254	254	254	253	253	253	164	162	161	161	163	Length
9	9	9	9	9	10	10	9	9	9	9	10	10	10	10	10	10	10	10	EG .
US-10-109-551-10	US-10-109-551-8	US-10-109-551-6	US-10-109-551-4	US-10-109-551-2	US-09-943-906-4	US-09-943-906-1	US-10-106-574-8	US-10-106-574-7	US-10-106-574-6	US-10-106-574-5	US-09-943-906-2	US-09-919-172-57	US-09-904-987-3	US-09-745-003-12	US-09-745-003-10	US-09-745-003-9	US-09-745-003-7	US-09-745-003-11	ID
Sequence 10, Appl	Sequence 8, Appli	Sequence 6, Appli	Sequence 4, Appli	Sequence 2, Appli	Sequence 4, Appli	Sequence 1, Appli	Sequence 8, Appli	Sequence 7, Appli	Sequence 6, Appli	Sequence 5, Appli	Sequence 2, Appli	Sequence 57, Appl	Sequence 3, Appli	Sequence 12, Appl	Sequence 10, Appl	Sequence 9, Appli	Sequence 7, Appli	Sequence 11, Appl	Description

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US-09-907-838-23 US-09-907-613-23 US-09-907-942-23 US-10-175-746-274	US-10-121-049-274 US-10-123-904-274 US-10-140-470-274	US-10-066-500-99 US-09-907-853-23 US-09-907-824-23 US-09-907-841-23 US-09-904-011-23 US-09-904-011-23	US-09-832-312-3 US-09-832-312-34 US-09-832-312-36 US-09-832-312-38 US-09-832-312-40 US-09-905-291A-23	US-09-943-906-3 US-10-209-194-2 US-10-115-984-2 US-09-832-312-7 US-09-832-312-9 US-09-832-312-9
Sequence 23, Appl Sequence 23, Appl Sequence 23, Appl Sequence 274, App	27223		Sequence 3, Appli Sequence 34, Appl Sequence 36, Appl Sequence 38, Appl Sequence 40, Appl Sequence 23, Appl	Sequence 3, Appli Sequence 2, Appli Sequence 2, Appli Sequence 7, Appli Sequence 9, Appli Sequence 5, Appli

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ALIGNMENTS

RESULT 1
US-09-745-003-11
US-09-745-003-11
; Sequence 11, Application US/09745003
; Patent No. US20020042122A1

GENERAL INFORMATION:
APPLICANT: Bazan, Fernando J
APPLICANTION: Human Proteins; Related Reagents

FILE REFERENCE: PrP2

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RESULT 2
US-09-745-003-7
Sequence 7, Application US/09745003
Patent No. US20020042122A1
Patent No. US20020042122A1
APPLICANT: Bazan, Fernando J
FILE OF INVENTION: Human Proteins; Related Reagents
FILE REFERENCE: PrP2
CURRENT APPLICATION NUMBER: US/09/745,003
CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION NUMBER: US/09/745,003;
CURRENT FILING DATE: 2000-12-20;
NUMBER OF SEQ ID NOS: 13;
SOFTWARE: PatentIn Ver. 2.0;
SEQ ID NO 11;
LEGIEN: 163;
TYPE: PRT
ORGANISM: Hamster sp.
                                                                                                                                                                                                                                                                                                        В
                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 8; Conserv
LENGTH: 161
TYPE: PRT
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RESULT 5
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US-09-745-003-10
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US-09-745-003-9
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GENERAL INFORMATION:
APPLICANT: Bazan, Fernando J
TITLE OF INVENTION: Human Proteins; Related Reagents
                                                                                                                                                                                                                                                                                                   SEQ ID NO 10
LENGTH: 162
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 10, Application US/09745003 Patent No. US20020042122A1 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       CURRENT APPLICATION NUMBER: US/09/745,003
CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO9
LENGTH: 161
TYPE: PRT
                                                 Sequence 12, Application US/09745003 Patent No. US20020042122A1
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Patent No. US20020042122A1
GENERAL INFORMATION:
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CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
                                                                                                                                                                                                                                                                                                                                                                                                            TITLE OF INVENTION: Human Proteins; Related Reagents FILE REFERENCE: prp2
                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: Bazan, Fernando J
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Local Similarity 61.5%;
les 8; Conserva++
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                                                                                                                                                                          1 CXTQYXXESXAXY 13
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                                                                                                                                             CITQYERESQAYY 135
                                                                                                                                                                                                         0; Mismatches
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Pred. No. 0.023;
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Pred. No. 0.023;
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US-09-904-987-3
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.US-09-745-003-12
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CURRENT FILING DATE: 2000-12-20
NUMBER OF SEQ ID NOS: 13
SOFTWARE: Patentin Ver. 2.0
SEQ ID NO 12
LENGTH: 164
TYPE: PRT
                                                                                                                                                                                                                                                                                                                                 Sequence 57, Application US/09919172 Patent No. US20020119463A1
                                                                                               SEQ ID NO 57
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Patent No. US20020037908A1
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Best Local Similarity 61.5%;
Matches 8; Conservative
                                                                                                                                                   CURRENT APPLICATION NUMBER: US/09/919,172
CURRENT FILING DATE: 2001-07-30
PRIOR APPLICATION NUMBER: 60/222,469
PRIOR FILING DATE: 2000-07-28
                                                                                                                                                                                                                            APPLICANT: Faris, Mary
APPLICANT: Turner, Christopher M.
TITLE OF INVENTION: PROSTATE CANCER MARKERS
FILE REFERENCE: PA-0036 US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LENGTH: 253

TYPE: PRT
ORGANISM: homo sapiens
ORGANISM: homo sapiens
PUBLICATION INFORMATION:
DATABASE ACCESSION NUMBER: NCBI ENTREZ / XM_009567
DATABASE ENTRY DATE: 2001-04-17
RELEVANT RESIDUES: (1)..(253)
                                                                                                                 NUMBER OF SEQ ID NOS: 102
SOFTWARE: PERL Program
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                CURRENT APPLICATION NUMBER: US/09/904,987
CURRENT FILING DATE: 2001-07-12
NUMBER OF SEQ ID NOS: 7
SOFTWARE: Patentin version 3.0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TITLE OF INVENTION: Methods and Compositions for Controlling Pathological and Prep
TITLE OF INVENTION: Protein Assembly or Aggregation
FILE REFERENCE: 42108/26146
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NAME/KEY: misc_feature
                    FEATURE:
                                  ORGANISM: Homo sapiens
                                                        LENGTH: 253
TYPE: PRT
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Pred. No. 0.023;
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; OTHER INFORMATION: Incyte ID No. US20020119463A1 1256895CD1 US-09-919-172-57
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US-09-943-906-2
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               US-10-106-574-5 
: Sequence 5, Application US/10106574 
: Patent No. US20020164335A1
                                                                                                                                                                                                                                                                       US-09-943-906-2
                                                                     RESULT 9
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GENERAL INFORMATION:
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INFORMATION FOR SEQ ID NO: 2:
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MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06
REFERENCE/DOCKET NUMBER: 06
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICATION NUMBER: US/09/943,906
FILING DATE: 30-Aug-2001
CLASSIFICATION: <Unknown>
PRIOR APPLICATION DATA:
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SOFTWARE: FBStSEQ Version 2.0
CURRENT APPLICATION DATA:
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Burton, Dennis R.
HITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE Pre
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         APPLICATION NUMBER: 09/550,374
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
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                                                                                                                                                                                                                                                                                    MOLECULE TYPE: peptide SEQUENCE DESCRIPTION: SEQ ID NO: 2:
                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear MOLECULE TYPE: peptid
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                                                                                                                                                                                                                                                                                                                                     LENGTH: 253 amino acids TYPE: amino acid STRANDEDNESS: single
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                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             NAME: Bozicevic, Karl
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Pred. No.
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RESULT 10
US-10-106-574-6
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                                                                      ; SOFTWARE: PatentIn version 3.1
; SEQ ID NO 7
; LENGTH: 254
; TYPE: PRT
; ORGANISM: Murinae gen. sp.
US-10-106-574-7
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US-10-106-574-7
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; SEQ ID NO 5
FNGTH: 254
                                                                                                                                                                               Sequence 7, Application US/10106574
Patent No. US20020164335A1
GENERAL INFORMATION:
APPLICANT: Harris, David A.
APPLICANT: Stewart, Richard S.
APPLICANT: Stewart, Richard S.
TITLE OF INVENTION: Compositions and Methods for the Study and Diagnosis of Prion
FILE REFERENCE: 09789280.0003
CURRENT APPLICATION NUMBER: US/10/106,574
CURRENT FILING DATE: 2002-03-26
NUMBER OF SEO ID NOS: 8
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LENGTH: 254
TYPE: PRT
ORGANISM: Murinae gen.
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Patent No. US20020164335A1
GENERAL INFORMATION:
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Matches 8; Conserv
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APPLICANT: Stewart, Richard S.

TITLE OF INVENTION: Compositions and Methods for the Study and Diagnosis of Prion FILE REFERENCE: 09789280.0003
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APPLICANT: Stewart, Richard S.
TITLE OF INVENTION: Compositions and Methods for the Study and Diagnosis of Prion
FILE REFERENCE: 09789280.0003
FILE REFERENCE: 09789280.0003
CURRENT APPLICATION NUMBER: US/10/106,574
CURRENT FILING DATE: 2002-03-26
CURRENT FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 8
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CURRENT FILLING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 8
SOFTWARE: PatentIn version 3.1
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Pred. No. 0.035;
0; Mismatches
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Pred. No. 0.035;
               Score 41; DB 9;
Pred. No. 0.035;
Mismatches
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US-09-943-906-1
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US-10-106-574-8
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Matches 8; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: Stewart, Richard S.
TITLE OF INVENTION: Compositions and Methods for the Study and Diagnosis of Prion Dis
FILE REFERENCE: 09789280.0003
CURRENT APPLICATION NUMBER: US/10/106,574
CURRENT FILING DATE: 2002-03-26
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CURRENT FILING DATE: 2002-03-26
NUMBER OF SEQ ID NOS: 8
SOFTWARE: Patentin version 3.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICANT: Harris, David A. APPLICANT: Stewart, Richar
                     TELEX: <Unknown>
INFORMATION FOR SEQ ID NO: 1:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         213 CVTOYOKESOAYY 225
                                                                                                                                                                                                                                                COMPUTER: IBM Compatible
OPERATING SYSTEM: DOS
SOFTWARE: FGASTEQ Version 2.0
CURRENT APPLICATION DATA:
APPLICATION UMBER: US/09/943,906
FILING DATE: 30-Aug-2001
CLASSIFICATION: CLASSIFICATION
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               213 CVTQYQKESQAYY 225
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                                                        NAME: BOZICEVIC, Karl
REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06510/059001
TELECOMMUNICATION IMFORMATION:
TELEPHONE: 415-854-5277
TELEFAX: 415-854-0875
                                                                                                                                                              APPLICATION NUMBER: 09/550,374
FILING DATE: <Unknown>
ATTORNEY/AGENT INFORMATION:
NAME: BOZICEVIC, Karl
                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Williamson, R. Anthony
Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE PrP
NUMBER OF SEQUENCES: 86
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       APPLICANT: Prusiner, Stanley B.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
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Pred. No. 0.035;
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US-09-943-906-4
; Sequence 4, Application US/09943906
; Patent No. US20020150571A1
; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
; SEQUENCE DESCRIPTION: SEQ ID NO: 1:
US-09-943-906-1
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Best Local
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INFORMATION FOR SEQ ID NO: 4:
SEQUENCE CHARACTERISTICS:
216 CITQYQRESQAYY 228
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                                                                            Local
                            1 CXTQYXXESXAXY 13
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                                                                                                                                         TOPOLOGY: linear
MOLECULE TYPE: peptide
SEQUENCE DESCRIPTION: SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                    PRIOR APPLICATION NUMBER: 09/550,374
APPLICATION NUMBER: 09/550,374
FILING DATE: <UNKNOWN>
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ZIP: 94025
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
                                                                                                                                                                                                                                                                                                      REGISTRATION NUMBER: 28,807
REFERENCE/DOCKET NUMBER: 06510/059001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 415-854-5277
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SOFTWARE: FastSEQ Version 2.0 CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NUMBER OF SEQUENCES: 80 CORRESPONDENCE ADDRESS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Williamson, R. Anthony
Burton, Dennis R.
TITLE OF INVENTION: ANTIBODIES SPECIFIC FOR NATIVE Prp
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               APPLICANT: Prusiner, Stanley B.
                                                            Similarity
8; Conserv
                                                                                                                                                                                      LENGTH: 255 amino acids TYPE: amino acid STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                      CLASSIFICATION: <Unknown>
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICATION NUMBER: US/09/943,906 FILING DATE: 30-Aug-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Com
OPERATING SYSTEM:
                                                                                                                                                                                                                                                                                          TELEFAX: 415-854-0875
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      STATE: CA
                                                                                                                                                                                                                                                                                                                                                                        NAME: Bozicevic,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Menlo Park
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STREET: 2200 Sand Hill Road
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ADDRESSEE: Fish & Richardson P.C.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           LENGTH: 254 amino acids
                                                              Conservative
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                                                                          80.4%;
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                                                                          Score 41; DB 10;
Pred. No. 0.036;
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                                                            Mismatches
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                                                                                         Length 255;
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RESULT 15
US-10-109-551-2
US-10-109-551-2
Sequence 2, Application US/10109551
Publication No. US20020194635A1
GENERAL INFORMATION:
APPLICANT: DIEDRAHITA, JORGE
FILLE OF INVENTION: TRANSGENIC ANIMALS RESISTANT TO TRANSMISSIBLE
FITILE OF INVENTION: SPONGIFORM ENCEPHALOPATHIES
FILLE OF INVENTION: SPONGIFORM ENCEPHALOPATHIES
FILLE REFERENCE: TAMK:207US
CURRENT FILLING DATE: 2002-03-28
PRIOR APPLICATION NUMBER: 05/10/109,551
CURRENT FILLING DATE: 2002-03-28
PRIOR FILLYGO DATE: 2001-03-30

NUMBER OF SEQ ID NOS: 10
SOFTWARE: Patentin Ver. 2.1
SEQ ID NO 2
LENGTH: 256
TYPE: PRT
CORGANISM: Bos taurus
US-10-109-551-2

Query Match
Best Local Similarity 61.5%; Pred. No. 0.036;
Matches 8: Conservative 0; Mismatches 5; Indels 0; Gaps
ON 1 CXTQYXXESXAXY 13
Db 217 CITQYQRESQAYY 229

Search completed: March 24, 2003, 17:25:05
Job time: 11.6458 secs
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